

Claims

1. A method for the detection of drink-spoiling microorganisms in a sample, whereby the detection is carried out by using at least one oligonucleotide probe  
5 having a nucleic acid sequence selected from the group consisting of (all sequences in 5' → 3'direction):

	SEQ ID No. 1:	5'- GTTGACCAAGATTCTCCGCTC
	SEQ ID No. 5:	5'- CCCGGTCGAATTAAAACC
10	SEQ ID No. 6:	5'- GCCCGGTCGAATTAAAAC
	SEQ ID No. 7:	5'- GGCCCGGTCGAATTAAAA
	SEQ ID No. 8:	5'- AGGCCCGGTCGAATTAAA
	SEQ ID No. 9:	5'- AAGGCCCGGTCGAATTAA
	SEQ ID No. 10:	5'- ATATTCGAGCGAACGCC
15	SEQ ID No. 11:	5'- AAAGATCCGGACCGGCCG
	SEQ ID No. 12	5'- GGAAAGATCCGGACCGGC
	SEQ ID No. 13	5'- GAAAGATCCGGACCGGCC
	SEQ ID No. 14	5'- GATCCGGACCGGCCGACC
	SEQ ID No. 15	5'- AGATCCGGACCGGCCGAC
20	SEQ ID No. 16	5'- AAGATCCGGACCGGCCGA
	SEQ ID No. 17	5'- GAAAGGCCCGGTCGAATT
	SEQ ID No. 18	5'- AAAGGCCCGGTCGAATT
	SEQ ID No. 19	5'- GGAAAGGCCCGGTCGAAT
	SEQ ID No. 20	5'- AGGAAAGGCCCGGTCGAA
25	SEQ ID No. 21	5'- AAGGAAAGGCCCGGTCGA
	SEQ ID No. 22:	5'- ATAGCACTGGGATCCTCGCC
	SEQ ID No. 23:	5'- CCAGCCCCAAAGTTACCTTC
	SEQ ID No. 24:	5'- TCCTTGACGTAAAGTCGCAG
	SEQ ID No. 25:	5'- GGAAGAAAACCAGTACGC
30	SEQ ID No. 26:	5'- CCGGTCGGAAGAAAACCA

	SEQ ID No. 27:	5'- GAAGAAAACCAGTACGCG
	SEQ ID No. 28:	5'- CCCGGTCGGAAGAAAAACC
	SEQ ID No. 29:	5'- CGGTCGGAAGAAAACCAG
	SEQ ID No. 30:	5'- GGTCGGAAGAAAACCAGT
5	SEQ ID No. 31:	5'- AAGAAAACCAGTACGCGG
	SEQ ID No. 32:	5'- GTACGCGGAAAAATCCGG
	SEQ ID No. 33:	5'- AGTACGCGGAAAAATCCG
	SEQ ID No. 34:	5'- GCGGAAAAATCCGGACCG
	SEQ ID No. 35:	5'- CGGAAGAAAACCAGTACG
10	SEQ ID No. 36:	5'- GCCCGGTCGGAAGAAAAC
	SEQ ID No. 37:	5'- CGCGGAAAAATCCGGACC
	SEQ ID No. 38:	5'- CAGTACGCGGAAAAATCC
	SEQ ID No. 39:	5'- AGAAAACCAGTACGCGGA
	SEQ ID No. 40:	5'- GGCCCCGGTCGGAAGAAAA
15	SEQ ID No. 41:	5'- ATAAACACCACCCGATCC
	SEQ ID No. 42:	5'- ACGCGGAAAAATCCGGAC
	SEQ ID No. 43:	5'- GAGAGGCCCGGTCGGAAG
	SEQ ID No. 44:	5'- AGAGGCCCGGTCGGAAGA
	SEQ ID No. 45:	5'- GAGGCCCGGTCGGAAGAA
20	SEQ ID No. 46:	5'- AGGCCCGGTCGGAAGAAA
	SEQ ID No. 47:	5'- CCGAGTGGGTCACTAAAT
	SEQ ID No. 48:	5'- CCAGTACGCGGAAAAATC
	SEQ ID No. 49:	5'- TAAACACCACCCGATCCC
	SEQ ID No. 50:	5'- GGAGAGGCCCGGTCGGAA
25	SEQ ID No. 51:	5'- GAAAACCAGTACGCGGAA
	SEQ ID No. 52:	5'- TACGCGGAAAAATCCGGA
	SEQ ID No. 53:	5'- GGCCACAGGGACCCAGGG
	SEQ ID No. 54:	5'- TCACCAAGGGCCACAGGG
	SEQ ID No. 55:	5'- GGGCCACAGGGACCCAGG
30	SEQ ID No. 56:	5'- TTCACCAAGGGCCACAGG

	SEQ ID No. 57:	5'- ACAGGGACCCAGGGCTAG
	SEQ ID No. 58:	5'- AGGGCCACAGGGACCCAG
	SEQ ID No. 59:	5'- GTTCACCAAGGGCCACAG
	SEQ ID No. 60:	5'- GCCACAGGGACCCAGGGC
5	SEQ ID No. 61:	5'- CAGGGACCCAGGGCTAGC
	SEQ ID No. 62:	5'- AGGGACCCAGGGCTAGCC
	SEQ ID No. 63:	5'- ACCAAGGGCCACAGGGAC
	SEQ ID No. 64:	5'- CCACAGGGACCCAGGGCT
	SEQ ID No. 65:	5'- CACAGGGACCCAGGGCTA
10	SEQ ID No. 66:	5'- CACCAAGGGCCACAGGGA
	SEQ ID No. 67:	5'- GGGACCCAGGGCTAGCCA
	SEQ ID No. 68:	5'- AGGAGAGGCCCGGTGGGA
	SEQ ID No. 69:	5'- AAGGAGAGGCCCGGTGG
	SEQ ID No. 70:	5'- GAAGGAGAGGCCCGGTG
15	SEQ ID No. 71:	5'- AGGGCTAGCCAGAAGGAG
	SEQ ID No. 72:	5'- GGGCTAGCCAGAAGGAGA
	SEQ ID No. 73:	5'- AGAAGGAGAGGCCCGGT
	SEQ ID No. 74:	5'- CAAGGGCCACAGGGACCC
	SEQ ID No. 75:	5'- CCAAGGGCCACAGGGACC
20	SEQ ID No. 76:	5'- GTCGGAAAAACCAGTACG
	SEQ ID No. 77:	5'- GCCCGGTCGGAAAAACCA
	SEQ ID No. 78:	5'- CCGGTGGAAAAACCA
	SEQ ID No. 79:	5'- CCCGGTCGGAAAAACCA
	SEQ ID No. 80:	5'- TCGGAAAAACCAGTACGC
25	SEQ ID No. 81:	5'- CGGAAAAACCAGTACGCG
	SEQ ID No. 82:	5'- GGAAAAACCAGTACGCGG
	SEQ ID No. 83:	5'- GTACGCGGAAAAATCCGG
	SEQ ID No. 84:	5'- AGTACGCGGAAAAATCCG
	SEQ ID No. 85:	5'- GCGGAAAAATCCGGACCG
30	SEQ ID No. 86:	5'- GGTGGAAAAACCAGTAC

	SEQ ID No. 87:	5'- ACTCCTAGTGGTGCCCTT
	SEQ ID No. 88:	5'- GCTCCACTCCTAGTGGTG
	SEQ ID No. 89:	5'- CACTCCTAGTGGTGCCCT
	SEQ ID No. 90:	5'- CTCCACTCCTAGTGGTGC
5	SEQ ID No. 91:	5'- TCCACTCCTAGTGGTGCC
	SEQ ID No. 92:	5'- CCACTCCTAGTGGTGCC
	SEQ ID No. 93:	5'- GGCTCCACTCCTAGTGGT
	SEQ ID No. 94:	5'- AGGCTCCACTCCTAGTGG
	SEQ ID No. 95:	5'- GGCCCCGGTCGGAAAAAACC
10	SEQ ID No. 96:	5'- GAAAAACCACTACGCGGA
	SEQ ID No. 97:	5'- CGCGGAAAAATCCGGACC
	SEQ ID No. 98:	5'- CAGTACGCGGAAAAATCC
	SEQ ID No. 99:	5'- CGGTCGGAAAAACCAGTA
	SEQ ID No. 100:	5'- AAGGCCCGGTCGGAAAAAA
15	SEQ ID No. 101:	5'- CAGGCTCCACTCCTAGTG
	SEQ ID No. 102:	5'- CTCCTAGTGGTGCCCTTC
	SEQ ID No. 103:	5'- TCCTAGTGGTGCCCTTCC
	SEQ ID No. 104:	5'- GCAGGCTCCACTCCTAGT
	SEQ ID No. 105:	5'- AGGCCCGGTCGGAAAAAC
20	SEQ ID No. 106:	5'- ACGCGGAAAAATCCGGAC
	SEQ ID No. 107:	5'- CCAGTACGCGGAAAAATC
	SEQ ID No. 108:	5'- CTAGTGGTGCCCTTCCGT
	SEQ ID No. 109:	5'- GAAAGGCCCGGTCGGAAA
	SEQ ID No. 110:	5'- AAAGGCCCGGTCGGAAAA
25	SEQ ID No. 111:	5'- TACGCGGAAAAATCCGGA
	SEQ ID No. 112:	5'- GGAAAGGCCCGGTCGGAA
	SEQ ID No. 113:	5'- ATCTCTTCCGAAAGGTGCG
	SEQ ID No. 114:	5'- CATCTCTTCCGAAAGGTGTC
	SEQ ID No. 115:	5'- CTCTTCCGAAAGGTGAGA
30	SEQ ID No. 116:	5'- CTTCCGAAAGGTGAGAT

	SEQ ID No. 117:	5'- TCTCTTCCGAAAGGTCGA
	SEQ ID No. 118:	5'- TCTTCCGAAAGGTCGAGA
	SEQ ID No. 119:	5'- CCTAGTGGTGCCCTTCCG
	SEQ ID No. 120:	5'- TAGTGGTGCCCTTCCGTC
5	SEQ ID No. 121:	5'- AGTGGTGCCCTTCCGTCA
	SEQ ID No. 122:	5'- GCCAAGGTTAGACTCGTT
	SEQ ID No. 123:	5'- GGCCAAGGTTAGACTCGT
	SEQ ID No. 124:	5'- CCAAGGTTAGACTCGTTG
	SEQ ID No. 125:	5'- CAAGGTTAGACTCGTTGG
10	SEQ ID No. 126:	5'- AAGGTTAGACTCGTTGGC
	SEQ ID No. 127:	5'- CTCGCCTCACGGGTTCTCA
	SEQ ID No. 128:	5'- GGCCC GGTCGAAATTAAA
	SEQ ID No. 129:	5'- AGGCCCGGTCGAAATTAA
	SEQ ID No. 130:	5'- AAGGCCCGGTCGAAATTAA
15	SEQ ID No. 131:	5'- AAAGGCCCGGTCGAAATT
	SEQ ID No. 132:	5'- GAAAGGCCCGGTCGAAAT
	SEQ ID No. 133:	5'- ATATTCGAGCGAAACGCC
	SEQ ID No. 134:	5'- GGAAAGGCCCGGTCGAAA
	SEQ ID No. 135:	5'- AAAGATCCGGACC GGCG
20	SEQ ID No. 136:	5'- GGAAAGATCCGGACC GGCG
	SEQ ID No. 137:	5'- GAAAGATCCGGACC GGCG
	SEQ ID No. 138:	5'- GATCCGGACC GGCGAEC
	SEQ ID No. 139:	5'- AGATCCGGACC GGCGAC
	SEQ ID No. 140:	5'- AAGATCCGGACC GGCGA
25	SEQ ID No. 141:	5'- AGGAAAGGCCCGGTCGAA
	SEQ ID No. 142:	5'- AAGGAAAGGCCCGGTCGA
	SEQ ID No. 143:	5'- CGAGCAAAACGCC TGCTTG
	SEQ ID No. 144:	5'- CGCTCTGAAAGAGAGTTGCC
	SEQ ID No. 145:	5'- AGTTGCCCTACACTAGAC..
30	SEQ ID No. 146:	5'- GCTTCTCCGTCCCGCGCCG

	SEQ ID No. 148:	5'- CCTGGTTCGCCAAAAAGGC
	SEQ ID No. 149:	5'-GATTCTCGGCCCATGGG
	SEQ ID No. 150:	5'- ACCCTCTACGGCAGCCTGTT
	SEQ ID No. 151:	5'- GATCGGTCTCCAGCGATTCA
5	SEQ ID No. 152:	5'- ACCCTCCACGGCGGCCTGTT
	SEQ ID No. 153:	5'- GATTCTCGCGCCATGGG
	SEQ ID No. 154:	5'- TCATCAGACGGGATTCTCAC
	SEQ ID No. 157:	5'-AGTTGCCCTCCTCTAAAGC
	SEQ ID No. 158:	5'-CTGCCACAAGGACAAATGGT
10	SEQ ID No. 159:	5'-TGCCCCCTCTCTAAGCAAAT
	SEQ ID No. 160:	5'-CCCCAAAGTTGCCCTCTC
	SEQ ID No. 163:	5'-AAGACCAGGCCACCTCAT
	SEQ ID No. 164:	5'- CATCATAGAACACCGTCC
	SEQ ID No. 165:	5'- CCTTCCGAAGTCGAGGTTTT
15	SEQ ID No. 166:	5'- GGGAGTGTGCAACTC
	SEQ ID No. 167:	5'- AGCGGTCGTTCGCAACCT
	SEQ ID No. 168:	5'- CCGAAGTCGGGTTTGC GG
	SEQ ID No. 169:	5'- GATAGCCAAACCAACCTTC
	SEQ ID No. 170:	5'- GCCGAAACCACCTTCAAAC
20	SEQ ID No. 171:	5'- GTGATAGCCAAACCAACCT
	SEQ ID No. 172:	5'- AGTGATAGCCAAACCAACCT
	SEQ ID No. 173:	5'- TTTAACGGATGCGTTCGAC
	SEQ ID No. 174:	5'- AAGTGATAGCCAAACCAACC
	SEQ ID No. 175:	5'- GGTTGAATACCGTCAACGTC
25	SEQ ID No. 176:	5'- GCACAGTATGTCAAGACCTG
	SEQ ID No. 177:	5'- CATCCGATGTGCAAGCACTT
	SEQ ID No. 178:	5'- TCATCCGATGTGCAAGCACT
	SEQ ID No. 179:	5'- CCGATGTGCAAGCACTTCAT
	SEQ ID No. 180:	5'- CCACTCATCCGATGTGCAAG
30	SEQ ID No. 181:	5'- GCCACAGTTCGCCACTCATC

	SEQ ID No. 182:	5'- CCTCCGCGTTGTCACCGGC
	SEQ ID No. 183:	5'- ACCAGTCGCCACAGTCGC
	SEQ ID No. 184:	5'- CACTCATCCGATGTGCAAGC
	SEQ ID No. 185:	5'- CCAGTCGCCACAGTCGCC
5	SEQ ID No. 186:	5'- CTCATCCGATGTGCAAGCAC
	SEQ ID No. 187:	5'- TCCGATGTGCAAGCACTTCA
	SEQ ID No. 188:	5'- CGCCACTCATCCGATGTGCA
	SEQ ID No. 189:	5'- CAGTCGCCACAGTCGCCA
	SEQ ID No. 190:	5'- GCCACTCATCCGATGTGCAA
10	SEQ ID No. 191:	5'- CGCCACAGTCGCCACTCAT
	SEQ ID No. 192:	5'- ATCCGATGTGCAAGCACTTC
	SEQ ID No. 193:	5'- GTTCGCCACAGTCGCCACT
	SEQ ID No. 194:	5'- TCCTCCGCGTTGTCACCGG
	SEQ ID No. 195:	5'- CGCCAGGGTTCATCCTGAGC
15	SEQ ID No. 196:	5'- AGTCGCCACAGTCGCCAC
	SEQ ID No. 197:	5'- TCGCCACAGTCGCCACTCA
	SEQ ID No. 198:	5'- TTAACGGGATGCGTTCGACT
	SEQ ID No. 199:	5'- TCGCCACTCATCCGATGTGC
	SEQ ID No. 200:	5'- CCACAGTCGCCACTCATCC
20	SEQ ID No. 201:	5'- GATTAAACGGGATGCGTTCG
	SEQ ID No. 202:	5'- TAACGGGATGCGTTCGACTT ---
	SEQ ID No. 203:	5'- AACGGGATGCGTTCGACTTG ---
	SEQ ID No. 204:	5'- CGAAGGTTACCGAACCGACT
	SEQ ID No. 205:	5'- CCGAAGGTTACCGAACCGAC
25	SEQ ID No. 206:	5'- CCCGAAGGTTACCGAACCGA
	SEQ ID No. 207:	5'- TTCCTCCGCGTTGTCACCG
	SEQ ID No. 208:	5'- CCGCCAGGGTTCATCCTGAG
	SEQ ID No. 209:	5'- TCCTTCCAGAAGTGATAGCC
	SEQ ID No. 210:	5'- CACCAGTCGCCACAGTCG
30	SEQ ID No. 211:	5'- ACGGGATGCGTTCGACTTGC ---

	SEQ ID No. 212:	5'- GTCCTTCCAGAAGTGATAGC
	SEQ ID No. 213:	5'- GCCAGGGTTCATCCTGAGCC
	SEQ ID No. 214:	5'- ACTCATCCGATGTGCAAGCA
	SEQ ID No. 215:	5'- ATCATTGCCTTGGTGAACCG
5	SEQ ID No. 216:	5'- TCCGCGTTGTCAACCGGCAG
	SEQ ID No. 217:	5'- TGAACCGTTACTCCACCAAC
	SEQ ID No. 218:	5'- GAAGTGATAGCCGAAACCAC
	SEQ ID No. 219:	5'- CCGCGTTGTCAACCGGCAGT
	SEQ ID No. 220:	5'- TTCGCCACTCATCCGATGTG
10	SEQ ID No. 221:	5'- CATTAAACGGGATGCGTTCG
	SEQ ID No. 222:	5'- CACAGTCGCCACTCATCCG
	SEQ ID No. 223:	5'- TTGCCACAGTCGCCACTC
	SEQ ID No. 224:	5'- CTCCGCGTTGTCAACCGGCA
	SEQ ID No. 225:	5'- ACGCCGCCAGGGTTCATCCT
15	SEQ ID No. 226:	5'- CCTTCCAGAAGTGATAAGCCG
	SEQ ID No. 227:	5'- TCATTGCCTTGGTGAACCGT
	SEQ ID No. 228:	5'- CACAGTATGTCAAGACCTGG
	SEQ ID No. 229:	5'- TTGGTGAACCGTTACTCCAC
	SEQ ID No. 230:	5'- CTTGGTGAACCGTTACTCCA
20	SEQ ID No. 231:	5'- GTGAACCGTTACTCCACCAA
	SEQ ID No. 232:	5'- GGCTCCCGAACGGTTACCGAA
	SEQ ID No. 233:	5'- GAAGGTTACCGAACCGAGFT
	SEQ ID No. 234:	5'- TGGCTCCCGAACGGTTACCGA
	SEQ ID No. 235:	5'- TAATACGCCGCCGGTCCTTC
25	SEQ ID No. 236:	5'- GAACCGTTACTCCACCAACT
	SEQ ID No. 237:	5'- TACGCCGCCGGTCCTTCAG
	SEQ ID No. 238:	5'- TCACCAGTTGCCACAGTTC
	SEQ ID No. 239:	5'- CCTTGGTGAACCGTTACTCC
	SEQ ID No. 240:	5'- CTCACCAGTTGCCACAGTT
30	SEQ ID No. 241:	5'- CGCCGCCAGGGTTCATCCTG

	SEQ ID No. 242:	5'- CCTTGGTGAACCATTACTCC
	SEQ ID No. 243:	5'- TGGTGAACCATTACTCCACC
	SEQ ID No. 244:	5'- GCCGCCAGGGTTCATCCTGA
	SEQ ID No. 245:	5'- GGTGAACCATTACTCCACCA
5	SEQ ID No. 246:	5'- CCAGGGTTCATCCTGAGCCA
	SEQ ID No. 247:	5'- AATACGCCGCCAGGGTCCTTCC
	SEQ ID No. 248:	5'- CACGCCGCCAGGGTTCATCC
	SEQ ID No. 249:	5'- AGTCGCCACTCATCCGATG
	SEQ ID No. 250:	5'- CGGGATGCGTTGACTTGCA
10	SEQ ID No. 251:	5'- CATTGCCTTGGTGAACCGTT
	SEQ ID No. 252:	5'- GCACGCCGCCAGGGTTCATC
	SEQ ID No. 253:	5'- CTTCCTCCGCCGTTGTCACC
	SEQ ID No. 254:	5'- TGGTGAACCGTTACTCCACC
	SEQ ID No. 255:	5'- CCTTCCTCCGCCGTTGTCAC-
15	SEQ ID No. 256:	5'- ACGCCGCCGGGTCCCTCCAGA
	SEQ ID No. 257:	5'- GGTGAACCGTTACTCCACCA
	SEQ ID No. 258:	5'- GGGTCCTCCAGAAGTGATA
	SEQ ID No. 259:	5'- CTTCCAGAAGTGATAGCCGA
	SEQ ID No. 260:	5'- GCCTTGGTGAACCATTACTC
20	SEQ ID No. 261:	5'- ACAGTCGCCACTCATCCGA
	SEQ ID No. 262:	5'- ACCTTCCTCCGCCGTTGTCAC
	SEQ ID No. 263:	5'- CGAACCGACTTGGGTGTTG
	SEQ ID No. 264:	5'- GAACCGACTTGGGTGTTGC
	SEQ ID No. 265:	5'- AGGTTACCGAACCGACTTTG
25	SEQ ID No. 266:	5'- ACCGAACCGACTTGGGTGT
	SEQ ID No. 267:	5'- TTACCGAACCGACTTGGGT
	SEQ ID No. 268:	5'- TACCGAACCGACTTGGGTG
	SEQ ID No. 269:	5'- GTTACCGAACCGACTTGGGG
	SEQ ID No. 270:	5'- CCTTCTGGTATGGTACCGTC
30	SEQ ID No. 271:	5'- TGCACCGCGGAYCCATCTCT

	SEQ ID No. 272:	5'- AGTTGCAGTCCAGTAAGCCG
	SEQ ID No. 273:	5'- GTTGCAGTCCAGTAAGCCGC
	SEQ ID No. 274:	5'- CAGTTGCAGTCCAGTAAGCC
	SEQ ID No. 275:	5'- TGCAGTCCAGTAAGCCGCCT
5	SEQ ID No. 276:	5'- TCAGTTGCAGTCCAGTAAGC
	SEQ ID No. 277:	5'- TTGCAGTCCAGTAAGCCGCC
	SEQ ID No. 278:	5'- GCAGTCCAGTAAGCCGCCCT
	SEQ ID No. 279:	5'- GTCAGTTGCAGTCCAGTAAG
	SEQ ID No. 280:	5'- CTCTAGGTGACGCCGAAGCG
10	SEQ ID No. 281:	5'- ATCTCTAGGTGACGCCGAAG
	SEQ ID No. 282:	5'- TCTAGGTGACGCCGAAGCGC
	SEQ ID No. 283:	5'- TCTCTAGGTGACGCCGAAGC
	SEQ ID No. 284:	5'- CCATCTCTAGGTGACGCCGA
	SEQ ID No. 285:	5'- CATCTCTAGGTGACGCCGAA
15	SEQ ID No. 286:	5'- TAGGTGACGCCGAAGCGCCT
	SEQ ID No. 287:	5'- CTAGGTGACGCCGAAGCGCC
	SEQ ID No. 288:	5'- CTTAGACGGCTCCTTCCTAA
	SEQ ID No. 289:	5'- CCTTAGACGGCTCCTTCCTA
	SEQ ID No. 290:	5'- ACGTCAGTTGCAGTCCAGTA
20	SEQ ID No. 291:	5'- CGTCAGTTGCAGTCCAGTAA
	SEQ ID No. 292:	5'- ACGCCGAAGCGCCTTTAAC
	SEQ ID No. 293:	5'- GACGCCGAAGCGCCTTTAA
	SEQ ID No. 294:	5'- GCCGAAGCGCCTTTAACTT
	SEQ ID No. 295:	5'- CGCCGAAGCGCCTTTAACT
25	SEQ ID No. 296:	5'- GTGACGCCGAAGCGCCTTT
	SEQ ID No. 297:	5'- TGACGCCGAAGCGCCTTTA
	SEQ ID No. 298:	5'- AGACGGCTCCTTCCTAAAAG
	SEQ ID No. 299:	5'- ACGGCTCCTTCCTAAAAGGT
	SEQ ID No. 300:	5'- GACGGCTCCTTCCTAAAAGG
30	SEQ ID No. 301:	5'- CCTTCCTAAAAGGTTAGGCC

	SEQ ID No. 302:	5'- GGTGACGCCAAAGCGCCTT
	SEQ ID No. 303:	5'- AGGTGACGCCAAAGCGCCTT
	SEQ ID No. 304:	5'- TAGGTGACGCCAAAGCGCCT
	SEQ ID No. 305:	5'- CTCTAGGTGACGCCAAAGCG
5	SEQ ID No. 306:	5'- TCTAGGTGACGCCAAAGCGC
	SEQ ID No. 307:	5'- CTAGGTGACGCCAAAGCGCC
	SEQ ID No. 308:	5'- ACGCCAAAGCGCCTTTAAC
	SEQ ID No. 309:	5'- CGCCAAAGCGCCTTTAACT
	SEQ ID No. 310:	5'- TGACGCCAAAGCGCCTTTA
10	SEQ ID No. 311:	5'- TCTCTAGGTGACGCCAAAGC
	SEQ ID No. 312:	5'- GTGACGCCAAAGCGCCTTT
	SEQ ID No. 313:	5'- GACGCCAAAGCGCCTTTAA
	SEQ ID No. 314:	5'- ATCTCTAGGTGACGCCAAAG
	SEQ ID No. 315:	5'- CATCTCTAGGTGACGCCAA
15	SEQ ID No. 316:	5'- TCCATCTCTAGGTGACGCCA
	SEQ ID No. 317:	5'- CCATCTCTAGGTGACGCCAA
	SEQ ID No. 318:	5'- CTGCCTTAGACGGCTCCCC
	SEQ ID No. 319:	5'- CCTGCCTTAGACGGCTCCCC
	SEQ ID No. 320:	5'- GTGTCATGCGACACTGAGTT
20	SEQ ID No. 321:	5'- TGTGTCATGCGACACTGAGT
	SEQ ID No. 322:	5'- CTTTGTGTCATGCGACACTG
	SEQ ID No. 323:	5'- TTGTGTCATGCGACACTGAG
	SEQ ID No. 324:	5'- TGCCTTAGACGGCTCCCCCT
	SEQ ID No. 325:	5'- AGACGGCTCCCCCTAAAAGG
25	SEQ ID No. 326:	5'- TAGACGGCTCCCCCTAAAAG
	SEQ ID No. 327:	5'- GCCTTAGACGGCTCCCCCTA
	SEQ ID No. 328:	5'- GCTCCCCCTAAAAGGTTAGG
	SEQ ID No. 329:	5'- GGCTCCCCCTAAAAGGTTAG
	SEQ ID No. 330:	5'- CTCCCCCTAAAAGGTTAGGC
30	SEQ ID No. 331:	5'- TCCCCCTAAAAGGTTAGGCC

	SEQ ID No. 332:	5'- CCCTAAAAGGTTAGGCCACC
	SEQ ID No. 333:	5'- CCCCTAAAAGGTTAGGCCAC
	SEQ ID No. 334:	5'- CGGCTCCCCCTAAAAGGTTA
	SEQ ID No. 335:	5'- CCCCCTAAAAGGTTAGGCCA
5	SEQ ID No. 336:	5'- CTTAGACGGCTCCCCCTAAA
	SEQ ID No. 337:	5'- TTAGACGGCTCCCCCTAAAA
	SEQ ID No. 338:	5'- GGGTCGCAACTCGTTGTAT
	SEQ ID No. 339:	5'- CCTTAGACGGCTCCCCCTAA
	SEQ ID No. 340:	5'- ACGGCTCCCCCTAAAAGGTT
10	SEQ ID No. 341:	5'- GACGGCTCCCCCTAAAAGGT
	SEQ ID No. 342:	5'- ACGCCGCAAGACCATCCTCT
	SEQ ID No. 343:	5'- CTAATACGCCGCAAGACCAT
	SEQ ID No. 344:	5'- TACGCCGCAAGACCATCCTC
	SEQ ID No. 345:	5'- GTTACGATCTAGCAAGCCGC
15	SEQ ID No. 346:	5'- AATAACGCCGCAAGACCATCC
	SEQ ID No. 347:	5'- CGCCGCAAGACCATCCTCTA
	SEQ ID No. 348:	5'- GCTAATACGCCGCAAGACCA
	SEQ ID No. 349:	5'- ACCATCCTCTAGCGATCCAA
	SEQ ID No. 350:	5'- TAATACGCCGCAAGACCATC
20	SEQ ID No. 351:	5'- AGCCATCCCTTTCTGGTAAG
	SEQ ID No. 352:	5'- ATACGCCGCAAGACCATCCT
	SEQ ID No. 353:	5'- AGTTACGATCTAGCAAGCCG
	SEQ ID No. 354:	5'- AGCTAATACGCCGCAAGACCC
	SEQ ID No. 355:	5'- GCCGCAAGACCATCCTCTAG
25	SEQ ID No. 356:	5'- TTACGATCTAGCAAGCCGCT
	SEQ ID No. 357:	5'- GACCATCCTCTAGCGATCCA
	SEQ ID No. 358:	5'- TTGCTACGTCACTAGGAGGC
	SEQ ID No. 359:	5'- ACGTCACTAGGAGGCGGAAA
	SEQ ID No. 360:	5'- TTTGCTACGTCACTAGGAGG
30	SEQ ID Nö. 361:	5'- GCCATCCCTTTCTGGTAAGG

	SEQ ID No. 362:	5'- TACGTCACTAGGAGGC GGAA
	SEQ ID No. 363:	5'- CGTCACTAGGAGGC GGAAAC
	SEQ ID No. 364:	5'- AAGACCATCCTCTAGCGATC
	SEQ ID No. 365:	5'- GCACGTATTAGCCATCCCT
5	SEQ ID No. 366:	5'- CTCTAGCGATCCAAAAGGAC
	SEQ ID No. 367:	5'- CCTCTAGCGATCCAAAAGGA
	SEQ ID No. 368:	5'- CCATCCTCTAGCGATCCAAA
	SEQ ID No. 369:	5'- GGCACGTATTAGCCATCCC
	SEQ ID No. 370:	5'- TACGATCTAGCAAGCCGCTT
10	SEQ ID No. 371:	5'- CAGTTACGATCTAGCAAGCC
	SEQ ID No. 372:	5'- CCGCAAGACCATCCTCTAGC
	SEQ ID No. 373:	5'- CCATCCCTTCTGGTAAGGT
	SEQ ID No. 374:	5'- AGACCATCCTCTAGCGATCC
	SEQ ID No. 375:	5'- CAAGACCATCCTCTAGCGAT
15	SEQ ID No. 376:	5'- GCTACGTCACTAGGAGGC GG
	SEQ ID No. 377:	5'- TGCTACGTCACTAGGAGGC G
	SEQ ID No. 378:	5'- CTACGTCACTAGGAGGC GGAA
	SEQ ID No. 379:	5'- CCTCAACGTCAGTTACGATC
	SEQ ID No. 380:	5'- GTCACTAGGAGGC GGAAACC
20	SEQ ID No. 381:	5'- TCCTCTAGCGATCCAAAAGG
	SEQ ID No. 382:	5'- TGGCACGTATTAGCCATCC
	SEQ ID No. 383:	5'- ACGATCTAGCAAGCCGCTTT
	SEQ ID No. 384:	5'- GCCAGTCTCTCAACTCGGCT
	SEQ ID No. 385:	5'- AAGCTAATACGCCGCAAGAC
25	SEQ ID No. 386:	5'- GTTGCTACGTCACTAGGAG
	SEQ ID No. 387:	5'- CGCCACTCTAGTCATTGCCT
	SEQ ID No. 388:	5'- GGCCAGCCAGTCTCTCAACT
	SEQ ID No. 389:	5'- CAGCCAGTCTCTCAACTCGG
	SEQ ID No. 390:	5'- CCCGAAGATCAATT CAGCGG
30	SEQ ID No. 391:	5'- CGGGCCAGTCTCTCAACTCG

	SEQ ID No. 392:	5'- CCAGCCAGTCTCTCAACTCG
	SEQ ID No. 393:	5'- TCATTGCCTCACTTCACCCG
	SEQ ID No. 394:	5'- GCCAGCCAGTCTCTCAACTC
	SEQ ID No. 395:	5'- CACCCGAAGATCAATT CAGC
5	SEQ ID No. 396:	5'- GTCATTGCCTCACTTCACCC
	SEQ ID No. 397:	5'- CATTGCCTCACTTCACCCGA
	SEQ ID No. 398:	5'- ATTGCCTCACTTCACCCGAA
	SEQ ID No. 399:	5'- CGAAGATCAATT CAGCGGCT
	SEQ ID No. 400:	5'- AGTCATTGCCTCACTTCACC
10	SEQ ID No. 401:	5'- TCGCCACTCTAGTCATTGCC
	SEQ ID No. 402:	5'- TTGCCTCACTTCACCCGAAG
	SEQ ID No. 403:	5'- CGGCCAGTCTCTCAACTCGG
	SEQ ID No. 404:	5'- CTGGCACGTATTAGCCATC
	SEQ ID No. 405:	5'- ACCCGAAGATCAATT CAGCG
15	SEQ ID No. 406:	5'- TCTAGCGATCCAAAAGGACC
	SEQ ID No. 407:	5'- CTAGCGATCCAAAAGGACCT
	SEQ ID No. 408:	5'- GCACCCATCGTTACGGTAT
	SEQ ID No. 409:	5'- CACCCATCGTTACGGTATG
	SEQ ID No. 410:	5'- GCCACTCTAGTCATTGCC
20	SEQ ID No. 411:	5'- CGTTGCTACGTCACTAGGA
	SEQ ID No. 412:	5'- GCCTCAACGTCAGTTACGAT
	SEQ ID No. 413:	5'- GCCGGCCAGTCTCTCAACTC
	SEQ ID No. 414:	5'- TCACTAGGAGGCGGAAACCT
	SEQ ID No. 415:	5'- AGCCTCAACGTCAGTTACGA
25	SEQ ID No. 416:	5'- AGCCAGTCTCTCAACTCGGC
	SEQ ID No. 417:	5'- GGCCAGTCTCTCAACTCGGC
	SEQ ID No. 418:	5'- CAAGCTAATACGCCGCAAGA
	SEQ ID No. 419:	5'- TTCGCCACTCTAGTCATTGC
	SEQ ID No. 420:	5'- CCGAAGATCAATT CAGCGGC
30	SEQ ID No. 421:	5'- CGCAAGACCATCCTCTAGCG

	SEQ ID No. 422:	5'- GCAAGACCATCCTCTAGCGA
	SEQ ID No. 423:	5'- GCGTTGCTACGTCACTAGG
	SEQ ID No. 424:	5'- CCACTCTAGTCATTGCCTCA
	SEQ ID No. 425:	5'- CACTCTAGTCATTGCCTCAC
5	SEQ ID No. 426:	5'- CCAGTCTCTCAACTCGGCTA
	SEQ ID No. 427:	5'- TTACCTTAGGCACCGGCCTC
	SEQ ID No. 428:	5'- ACAAGCTAATACGCCGCAAG
	SEQ ID No. 429:	5'- TTTACCTTAGGCACCGGCCT
	SEQ ID No. 430:	5'- TTTTACCTTAGGCACCGGCC
10	SEQ ID No. 431:	5'- ATTTTACCTTAGGCACCGGC
	SEQ ID No. 432:	5'- GATTTCACCTTAGGCACCGG
	SEQ ID No. 433:	5'- CTCACTTCACCCGAAGATCA
	SEQ ID No. 434:	5'- ACGCCACCAAGCGTTCATCCT
	SEQ ID No. 435:	5'- GCCAAGCGACTTGGGTACT
15	SEQ ID No. 436:	5'- CGGAAAATTCCCTACTGCAG
	SEQ ID No. 437:	5'- CGATCTAGCAAGCCGCTTTC
	SEQ ID No. 438:	5'- GGTACCGTCAAGCTGAAAAC
	SEQ ID No. 439:	5'- TGCCTCACCCACCCGAAGA
	SEQ ID No. 440:	5'- GGCCGGCCAGTCTCTCAACT
20	SEQ ID No. 441:	5'- GGTAAGGTACCGTCAAGCTG
	SEQ ID No. 442:	5'- GTAAGGTACCGTCAAGCTGA
	SEQ ID No. 443:	5'- CCGCAAGACCATCCTCTAGG
	SEQ ID No. 444:	5'- ATTAGCCATCCCTTTCTGG
	SEQ ID No. 445:	5'- AACCCCTTCATCACACACCG
25	SEQ ID No. 446:	5'- CGAAACCCCTTCATCACAC
	SEQ ID No. 447:	5'- ACCCTTCATCACACACACGC
	SEQ ID No. 448:	5'- TACCGTCACACACTGAAC
	SEQ ID No. 449:	5'- AGATACCGTCACACACTG
	SEQ ID No. 450:	5'- CACTCAAGGGCGGAAACC
30	SEQ ID No. 451:	5'- ACCGTCACACACTGAACA

	SEQ ID No. 452:	5'- CGTCACACACTGAACAGT
	SEQ ID No. 453:	5'- CCGAAACCCTTCATCACA
	SEQ ID No. 454:	5'- CCGTCACACACTGAACAG
	SEQ ID No. 455:	5'- GATACCGTCACACACTGA
5	SEQ ID No. 456:	5'- GGTAAGATAACCGTCACAC
	SEQ ID No. 457:	5'- CCCTTCATCACACACGCG
	SEQ ID No. 458:	5'- ACAGTGTTCACGAGCCG
	SEQ ID No. 459:	5'- CAGTGTTCACGAGCCGA
	SEQ ID No. 460:	5'- ACAAAGCGTTCGACTTGC
10	SEQ ID No. 461:	5'- CGGATAACGCTTGGAACAA
	SEQ ID No. 462:	5'- AGGGCGGAAACCCCTCGAA
	SEQ ID No. 463:	5'- GGGCGGAAACCCCTCGAAC
	SEQ ID No. 464:	5'- GGCGGAAACCCCTCGAACAA
	SEQ ID No. 465:	5'- TGAGGGCTTCACTTCACTCAG
15	SEQ ID No. 466:	5'- AGGGCTTCACTTCACTCAGAC
	SEQ ID No. 467:	5'- GAGGGCTTCACTTCACTCAGA
	SEQ ID No. 468:	5'- ACTGCACTCAAGTCATCC
	SEQ ID No. 469:	5'- CCGGATAACGCTTGGAAC
	SEQ ID No. 470:	5'- TCCGGATAACGCTTGGAA
20	SEQ ID No. 471:	5'- TATCCCCTGCTAAGAGGT
	SEQ ID No. 472:	5'- CCTGCTAAGAGGTAGGTT
	SEQ ID No. 473:	5'- CCCTGCTAAGAGGTAGGT
	SEQ ID No. 474:	5'- CCCCTGCTAAGAGGTAGG
	SEQ ID No. 475:	5'- TCCCCTGCTAAGAGGTAG
25	SEQ ID No. 476:	5'- ATCCCCTGCTAAGAGGTA
	SEQ ID No. 477:	5'- CCGTTCCCTTCTGGTAAG
	SEQ ID No. 478:	5'- GCCGTTCCCTTCTGGTAA
	SEQ ID No. 479:	5'- AGCCGTTCCCTTCTGGTA
	SEQ ID No. 480:	5'- GCACGTATTAGCCGTTCC
30	SEQ ID No. 481:	5'- CACGTATTAGCCGTTCC

	SEQ ID No. 482:	5'- GGCACGTATTAAGCCGTT
	SEQ ID No. 483:	5'- CACTTCCCTCTACTGCAC
	SEQ ID No. 484:	5'- CCACTTCCCTCTACTGCA
	SEQ ID No. 485:	5'- TCCACTTCCCTCTACTGC
5	SEQ ID No. 486:	5'- CTTCCCTCTACTGCACTC
	SEQ ID No. 487:	5'- TAGCCGTTCCCTTCTGGT
	SEQ ID No. 488:	5'- TTAGCCGTTCCCTTCTGG
	SEQ ID No. 489:	5'- TTATCCCCTGCTAAGAGG
	SEQ ID No. 490:	5'- GTTATCCCCTGCTAAGAG
10	SEQ ID No. 491:	5'- CCCGTTGCCACTCTTG
	SEQ ID No. 492:	5'- AGCTGAGGGCTTCACTT
	SEQ ID No. 493:	5'- GAGCTGAGGGCTTCACT
	SEQ ID No. 494:	5'- GCTGAGGGCTTCACTTC
	SEQ ID No. 495:	5'- CTGAGGGCTTCACTTCA
15	SEQ ID No. 496:	5' CCCGTGTCCCGAAGGAAC
	SEQ ID No. 497:	5' GCACGAGTATGTCAAGAC
	SEQ ID No. 498:	5' GTATCCCGTGTCCCGAAG
	SEQ ID No. 499:	5' TCCCGTGTCCCGAAGGAA
	SEQ ID No. 500:	5' ATCCCGTGTCCCGAAGGA
20	SEQ ID No. 501:	5' TATCCCGTGTCCCGAAGG
	SEQ ID No. 502:	5' CTTACCTTAGGAAGCGCC
	SEQ ID No. 503:	5' TTACCTTAGGAAGCGCCC
	SEQ ID No. 504:	5' CCTGTATCCCGTGTCCCG
	SEQ ID No. 505:	5' CCACCTGTATCCCGTGT
25	SEQ ID No. 506:	5' CACCTGTATCCCGTGTCC
	SEQ ID No. 507:	5' ACCTGTATCCCGTGTCCC
	SEQ ID No. 508:	5' CTGTATCCCGTGTCCCGA
	SEQ ID No. 509:	5' TGTATCCCGTGTCCCGAA
	SEQ ID No. 510:	5' CACGAGTATGTCAAGACC
30	SEQ ID No. 511:	5' CGGTCTTACCTTAGGAAG

	SEQ ID No. 512:	5' TAGGAAGGCCCTCCTG
	SEQ ID No. 513:	5' AGGAAGGCCCTCCTTGC
	SEQ ID No. 514:	5' TTAGGAAGGCCCTCCTT
	SEQ ID No. 515:	5' CTTAGGAAGGCCCTCCT
5	SEQ ID No. 516:	5' CCTTAGGAAGGCCCTCC
	SEQ ID No. 517:	5' ACCTTAGGAAGGCCCTC
	SEQ ID No. 518:	5' TGCACACAATGGTGAGC
	SEQ ID No. 519:	5' TACCTTAGGAAGGCCCT
	SEQ ID No. 520:	5' ACCACCTGTATCCGTGT
10	SEQ ID No. 521:	5' GCACCACCTGTATCCCGT
	SEQ ID No. 522:	5' CACCACCTGTATCCGTG
	SEQ ID No. 523:	5' GCGGTTAGGCAACCTACT
	SEQ ID No. 524:	5' TGCAGGTTAGGCAACCTAC
	SEQ ID No. 525:	5' TTGCGGTTAGGCAACCTA
15	SEQ ID No. 526:	5' GGTCTTACCTTAGGAAGC
	SEQ ID No. 527:	5' GCTAATACAACGCGGGAT
	SEQ ID No. 528:	5' CTAATACAACGCGGGATC
	SEQ ID No. 529:	5' ATACAACGCGGGATCATC
	SEQ ID No. 530:	5' CGGTTAGGCAACCTACTT
20	SEQ ID No. 531:	5' TGCACCAACCTGTATCCCG
	SEQ ID No. 532:	5' GAAGGCCCTCCTGCGG
	SEQ ID No. 533:	5' GGAAGGCCCTCCTGCG
	SEQ ID No. 534:	5' CGTCCCTTCTGGTTAGA
	SEQ ID No. 535:	5' AGCTAATACAACGCGGG
25	SEQ ID No. 536:	5' TAGCTAATACAACGCGGG
	SEQ ID No. 537:	5' CTAGCTAATACAACGCGG
	SEQ ID No. 538:	5' GGCTATGTATCATGCCT
	SEQ ID No. 539:	5' GAGCCACTGCCTTTACA
	SEQ ID No. 540:	5' GTCGGCTATGTATCATCG
30	SEQ ID No. 541:	5' GGTGGCTATGTATCATC

	SEQ ID No. 542:	5' CAGGTCGGCTATGTATCA
	SEQ ID No. 543:	5' CGGCTATGTATCATCGCC
	SEQ ID No. 544:	5' TCGGCTATGTATCATCGC
	SEQ ID No. 545:	5' GTCTTACCTTAGGAAGCG
5	SEQ ID No. 546:	5' TCTTACCTTAGGAAGCGC
	SEQ ID No. 547:	5'- GTACAAACCGCCTACACGCC
	SEQ ID No. 548:	5'- TGTACAAACCGCCTACACGC
	SEQ ID No. 549:	5'- GATCAGCACGATGTCGCCAT
	SEQ ID No. 550:	5'- CTGTACAAACCGCCTACACG
10	SEQ ID No. 551:	5'- GAGATCAGCACGATGTCGCC
	SEQ ID No. 552:	5'- AGATCAGCACGATGTCGCCA
	SEQ ID No. 553:	5'- ATCAGCACGATGTCGCCATC
	SEQ ID No. 554:	5'- TCAGCACGATGTCGCCATCT
	SEQ ID No. 555:	5'- ACTGTACAAACCGCCTACAC
15	SEQ ID No. 556:	5'- CCGCCACTAAGGCCGAAACC
	SEQ ID No. 557:	5'- CAGCACGATGTCGCCATCTA
	SEQ ID No. 558:	5'- TACAAACCGCCTACACGCC
	SEQ ID No. 559:	5'- AGCACGATGTCGCCATCTAG
	SEQ ID No. 560:	5'- CGGCTTTAGAGATCAGCAC
20	SEQ ID No. 561:	5'- TCCGCCACTAAGGCCGAAAC
	SEQ ID No. 562:	5'- GACTGTACAAACCGCCTACA
	SEQ ID No. 563:	5'- GTCCGCCACTAAGGCCGAAA
	SEQ ID No. 564:	5'- GGGGATTTCACATCTGACTG
	SEQ ID No. 565:	5'- CATACAAGCCCTGGTAAGGT
25	SEQ ID No. 566:	5'- ACAAGCCCTGGTAAGGTTCT
	SEQ ID No. 567:	5'- ACAAAACCGCCTACACGCCCT
	SEQ ID No. 568:	5'- CTGACTGTACAAACCGCCTA
	SEQ ID No. 569:	5'- TGACTGTACAAACCGCCTAC
	SEQ ID No. 570:	5'- ACGATGTCGCCATCTAGCTT
30	SEQ ID No. 571:	5'- CACGATGTCGCCATCTAGCT

	SEQ ID No. 572:	5'- CGATGTCGCCATCTAGCTTC
	SEQ ID No. 573:	5'- GCACGATGTCGCCATCTAGC
	SEQ ID No. 574:	5'- GATGTCGCCATCTAGCTTCC
	SEQ ID No. 575:	5'- ATGTCGCCATCTAGCTTCCC
5	SEQ ID No. 576:	5'- TGTCGCCATCTAGCTTCCA
	SEQ ID No. 577:	5'- GCCATCTAGCTTCCCACTGT
	SEQ ID No. 578:	5'- TCGCCATCTAGCTTCCCACT
	SEQ ID No. 579:	5'- CGCCATCTAGCTTCCCACTG
	SEQ ID No. 580:	5'- GTCGCCATCTAGCTTCCAC
10	SEQ ID No. 581:	5'- TACAAGCCCTGGTAAGGTTTC
	SEQ ID No. 582:	5'- GCCACTAAGGCCGAAACCTT
	SEQ ID No. 583:	5'- ACTAAGGCCGAAACCTTCGT
	SEQ ID No. 584:	5'- CTAAGGCCGAAACCTTCGTG
	SEQ ID No. 585:	5'- CACTAAGGCCGAAACCTTCG
15	SEQ ID No. 586:	5'- AAGGCCGAAACCTTCGTGCG
	SEQ ID No. 587:	5'- CCACTAAGGCCGAAACCTTC
	SEQ ID No. 588:	5'- TAAGGCCGAAACCTTCGTGC
	SEQ ID No. 589:	5'- AGGCCGAAACCTTCGTGCGA
	SEQ ID No. 590:	5'- TCTGACTGTACAAACCGCCT
20	SEQ ID No. 591:	5'- CATCTGACTGTACAAACCGC
	SEQ ID No. 592:	5'- ATCTGACTGTACAAACCGCC
	SEQ ID No. 593:	5'- GTTCGTGCGACTTGCATGTG
	SEQ ID No. 594:	5'- CCTTCGTGCGACTTGCATGT
	SEQ ID No. 595:	5'- CTCTCTAGAGTGCCCCACCCA
25	SEQ ID No. 596:	5'- TCTCTAGAGTGCCCCACCAA
	SEQ ID No. 597:	5'- ACGTATCAAATGCAGCTCCC
	SEQ ID No. 598:	5'- CGTATCAAATGCAGCTCCC
	SEQ ID No. 599:	5'- CGCCACTAAGGCCGAAACCT
	SEQ ID No. 600:	5'- CCGAACCTTCGTGCGACTT
30	SEQ ID No. 601:	5'- GCCGAAACCTTCGTGCGACTT

	SEQ ID No. 602:	5'- AACCTTCGTGCGACTTGCAT
	SEQ ID No. 603:	5'- CGAACCTTCGTGCGACTTG
	SEQ ID No. 604:	5'- ACCTTCGTGCGACTTGCATG
	SEQ ID No. 605:	5'- GAAACCTTCGTGCGACTTGC
5	SEQ ID No. 606:	5'- GGCGAACCTTCGTGCGAC
	SEQ ID No. 607:	5'- AAACCTTCGTGCGACTTGCA
	SEQ ID No. 608:	5'- CACGTATCAAATGCAGCTCC
	SEQ ID No. 609:	5'- GCTCACCGGCTTAAGGTCAA
	SEQ ID No. 610:	5'- CGCTCACCGGCTTAAGGTCA
10	SEQ ID No. 611:	5'- TCGCTCACCGGCTTAAGGTCA
	SEQ ID No. 612:	5'- CTCACCGGCTTAAGGTCAA
	SEQ ID No. 613:	5'- CCCGACCGTGGTCGGCTGCG
	SEQ ID No. 614:	5'- GCTCACCGGCTTAAGGTCAA
	SEQ ID No. 615:	5'- CGCTCACCGGCTTAAGGTCA
15	SEQ ID No. 616:	5'- TCGCTCACCGGCTTAAGGTCA
	SEQ ID No. 617:	5'- CTCACCGGCTTAAGGTCAA
	SEQ ID No. 618:	5'- CCCGACCGTGGTCGGCTGCG
	SEQ ID No. 619:	5'- TCACCGGCTTAAGGTCAAAC
	SEQ ID No. 620:	5'- CAACCCTCTCTCACACTCTA
20	SEQ ID No. 621:	5'- ACAACCCTCTCTCACACTCT
	SEQ ID No. 622:	5'- CCACAACCCTCTCTCACACT
	SEQ ID No. 623:	5'- AACCCTCTCTCACACTCTAG
	SEQ ID No. 624:	5'- CACAACCCTCTCTCACACTC
	SEQ ID No. 625:	5'- TCCACAACCCTCTCTCACAC
25	SEQ ID No. 626:	5'- TTCCACAACCCTCTCTCAC
	SEQ ID No. 627:	5'- ACCCTCTCTCACACTCTAGT
	SEQ ID No. 628:	5'- GAGCCAGGTTGCCGCCTCG
	SEQ ID No. 629:	5'- AGGTCAAACCAACTCCCAGT
	SEQ ID No. 630:	5'- ATGAGCCAGGTTGCCGCCTT
30	SEQ ID No. 631:	5'- TGAGCCAGGTTGCCGCCTTC

	SEQ ID No. 632:	5'- AGGCTCCTCCACAGGCGACT
	SEQ ID No. 633:	5'- CAGGCTCCTCCACAGGCGAC
	SEQ ID No. 634:	5'- GCAGGCTCCTCCACAGGCGA
	SEQ ID No. 635:	5'- TTCGCTCACCGGCTTAAGGT
5	SEQ ID No. 636:	5'- GTTCGCTCACCGGCTTAAGG
	SEQ ID No. 637:	5'- GGTCGCTCACCGGCTTAAG
	SEQ ID No. 638:	5'- ATTCCACAACCCTCTCAC
	SEQ ID No. 639:	5'- TGACCCGACCGTGGTCGGCT
	SEQ ID No. 640:	5'- CCCTCTCTCACACTCTAGTC
10	SEQ ID No. 641:	5'- GAATTCCACAACCCTCTCTC
	SEQ ID No. 642:	5'- AGCCAGGTTGCCGCCTTCGC
	SEQ ID No. 643:	5'- GCCAGGTTGCCGCCTTCGCC
	SEQ ID No. 644:	5'- GGAATTCCACAACCCTCTCT
	SEQ ID No. 645:	5'- GGGATTCCACAACCCTCTC
15	SEQ ID No. 646:	5'- AACGCAGGCTCCTCCACAGG
	SEQ ID No. 647:	5'- CGGCTTAAGGTCAAACCAAC
	SEQ ID No. 648:	5'- CCGGCTTAAGGTCAAACCAA
	SEQ ID No. 649:	5'- CACCGGCTTAAGGTCAAACC
	SEQ ID No. 650:	5'- ACCGGCTTAAGGTCAAACCA
20	SEQ ID No. 651:	5'- ACCAACATCCAGCACACAT
	SEQ ID No. 652:	5'- TCGCTGACCCGACCGTGGTC
	SEQ ID No. 653:	5'- CGCTGACCCGACCGTGGTCG
	SEQ ID No. 654:	5'- GACCCGACCGTGGTCGGCTG
	SEQ ID No. 655:	5'- GCTGACCCGACCGTGGTCGG
25	SEQ ID No. 656:	5'- CTGACCCGACCGTGGTCGGC
	SEQ ID No. 657:	5'- CAGGCAGCTTGCCTTGA
	SEQ ID No. 658:	5'- TCATGCGGTATTAGCTCCAG
	SEQ ID No. 659:	5'- ACTAGCTAATCGAACGCCAGG
	SEQ ID No. 660:	5'- CATGCGGTATTAGCTCCAGT
30	SEQ ID No. 661:	5'- CGCAGGCTCCTCCACAGGCG

	SEQ ID No. 662:	5' - ACGCAGGCTCCTCCACAGGC
	SEQ ID No. 663:	5' - CTCAGGTGTCATGCGGTATT
	SEQ ID No. 664:	5' - CGCCTTGACCCTCAGGTGT
	SEQ ID No. 665:	5' - ACCCTCAGGTGTCATGCGGT
5	SEQ ID No. 666:	5' - CCTCAGGTGTCATGCGGTAT
	SEQ ID No. 667:	5' - TTTGACCCTCAGGTGTCATG
	SEQ ID No. 668:	5' - GACCCTCAGGTGTCATGCGG
	SEQ ID No. 669:	5' - TGACCCTCAGGTGTCATGCG
	SEQ ID No. 670:	5' - GCCTTGACCCTCAGGTGTC
10	SEQ ID No. 671:	5' - TTGACCCTCAGGTGTCATGC
	SEQ ID No. 672:	5' - CCCTCAGGTGTCATGCGGT
	SEQ ID No. 673:	5' - CCTTGACCCTCAGGTGTCA
	SEQ ID No. 674:	5' - CTTTGACCCTCAGGTGTCAT
	SEQ ID No. 675:	5' - AGTTATCCCCACCCATGGA
15	SEQ ID No. 676:	5' - CCAGCTATCGATCATCGCCT
	SEQ ID No. 677:	5' - ACCAGCTATCGATCATCGCC
	SEQ ID No. 678:	5' - CAGCTATCGATCATCGCCTT
	SEQ ID No. 679:	5' - AGCTATCGATCATCGCCTTG
	SEQ ID No. 680:	5' - GCTATCGATCATCGCCTTGG
20	SEQ ID No. 681:	5' - CTATCGATCATCGCCTTGGT
	SEQ ID No. 682:	5' - TTCGTGCGACTTGCATGTGT
	SEQ ID No. 683:	5' - TCGATCATCGCCTTGGTAGG
	SEQ ID No. 684:	5' - ATCGATCATCGCCTTGGTAG
	SEQ ID No. 685:	5' - CACAGGCGACTTGCAGCCTT
25	SEQ ID No. 686:	5' - CCACAGGCGACTTGCAGCCTT
	SEQ ID No. 687:	5' - TCCACAGGCGACTTGCAGCCT
	SEQ ID No. 688:	5' - TCCTCCACAGGCGACTTGCAG
	SEQ ID No. 689:	5' - CCTCCACAGGCGACTTGCAGC
	SEQ ID No. 690:	5' - CTCCACAGGCGACTTGCAGC
30	SEQ ID No. 691:	5' - ACAGGCGACTTGCAGCCTTGG

	SEQ ID No. 692:	5'- GCTCACCGGCTTAAGGTCAA
	SEQ ID No. 693:	5'- CGCTCACCGGCTTAAGGTCA
	SEQ ID No. 694:	5'- TCGCTCACCGGCTTAAGGTC
	SEQ ID No. 695:	5'- CTCACCGGCTTAAGGTCAAA
5	SEQ ID No. 696:	5'- CCCGACCGTGGTCGGCTGCG
	SEQ ID No. 697:	5'- TCACCGGCTTAAGGTCAAAC
	SEQ ID No. 698:	5'- CAACCCTCTCTCACACTCTA
	SEQ ID No. 699:	5'- ACAACCCTCTCTCACACTCT
	SEQ ID No. 700:	5'- CCACAACCCTCTCTCACACT
10	SEQ ID No. 701:	5'- AACCCCTCTCTCACACTCTAG
	SEQ ID No. 702:	5'- CACAACCCTCTCTCACACTC
	SEQ ID No. 703:	5'- TCCACAACCCTCTCTCACAC
	SEQ ID No. 704:	5'- TTCCACAACCCTCTCTCACA
	SEQ ID No. 705:	5'- ACCCTCTCTCACACTCTAGT
15	SEQ ID No. 706:	5'- GAGCCAGGTTGCCGCCTCG
	SEQ ID No. 707:	5'- AGGTCAAACCAACTCCCATG
	SEQ ID No. 708:	5'- ATGAGCCAGGTTGCCGCCTT
	SEQ ID No. 709:	5'- TGAGCCAGGTTGCCGCCTTC
	SEQ ID No. 710:	5'- AGGCTCCTCCACAGGCGACT
20	SEQ ID No. 711:	5'- CAGGCTCCTCCACAGGCGAC
	SEQ ID No. 712:	5'- GCAGGCTCCTCCACAGGCGA
	SEQ ID No. 713:	5'- TTTCGCTCACCGGCTTAAGGT
	SEQ ID No. 714:	5'- GTTCGCTCACCGGCTTAAGG
	SEQ ID No. 715:	5'- GGTCGCTCACCGGCTTAAG
25	SEQ ID No. 716:	5'- ATTCCACAACCCTCTCAC
	SEQ ID No. 717:	5'- TGACCCGACCGTGGTCGGCT
	SEQ ID No. 718:	5'- CCCTCTCTCACACTCTAGTC
	SEQ ID No. 719:	5'- GAATTCCACAACCCTCTCTC
	SEQ ID No. 720:	5'- AGCCAGGTTGCCGCCTCGC
30	SEQ ID No. 721:	5'- GCCAGGTTGCCGCCTCGCC

	SEQ ID No. 722:	5'- GGAATTCCACAACCCCTCT
	SEQ ID No. 723:	5'- GGGATTCCACAACCCCTCTC
	SEQ ID No. 724:	5'- AACGCAGGCTCCTCCACAGG
	SEQ ID No. 725:	5'- CGGCTTAAGGTCAAACCAAC
5	SEQ ID No. 726:	5'- CCGGCTTAAGGTCAAACCAA
	SEQ ID No. 727:	5'- CACCGGCTTAAGGTCAAACC
	SEQ ID No. 728:	5'- ACCGGCTTAAGGTCAAACCA
	SEQ ID No. 729:	5'- ACCAACATCCAGCACACAT
	SEQ ID No. 730:	5'- TCGCTGACCCGACCGTGGTC
10	SEQ ID No. 731:	5'- CGCTGACCCGACCGTGGTCG
	SEQ ID No. 732:	5'- GACCCGACCGTGGTCGGCTG
	SEQ ID No. 733:	5'- GCTGACCCGACCGTGGTCGG
	SEQ ID No. 734:	5'- CTGACCCGACCGTGGTCGGC
	SEQ ID No. 735:	5'- CAGGCGACTTGCGCCTTGA
15	SEQ ID No. 736:	5'- TCATCGGGTATTAGCTCCAG
	SEQ ID No. 737:	5'- ACTAGCTAATCGAACGCAGG
	SEQ ID No. 738:	5'- CATCGGGTATTAGCTCCAGT
	SEQ ID No. 739:	5'- CGCAGGCTCCTCCACAGGCG
	SEQ ID No. 740:	5'- ACGCAGGCTCCTCCACAGGC
20	SEQ ID No. 741:	5'- CTCAGGTGTCATGCGGTATT
	SEQ ID No. 742:	5'- CGCCTTGACCCCTCAGGTGT
	SEQ ID No. 743:	5'- ACCCTCAGGTGTATGCGGT
	SEQ ID No. 744:	5'- CCTCAGGTGTCATGCGGTAT
	SEQ ID No. 745:	5'- TTTGACCCCTCAGGTGTATG
25	SEQ ID No. 746:	5'- GACCCTCAGGTGTATGCGG
	SEQ ID No. 747:	5'- TGACCCTCAGGTGTATGCG
	SEQ ID No. 748:	5'- GCCTTGACCCCTCAGGTGTC
	SEQ ID No. 749:	5'- TTGACCCCTCAGGTGTATGC
	SEQ ID No. 750:	5'- CCCTCAGGTGTATGCGGTA
30	SEQ ID No. 751:	5'- CCTTGACCCCTCAGGTGTCA

	SEQ ID No. 752:	5'- CTTGACCCTCAGGTGTCAT
	SEQ ID No. 753:	5'- AGTTATCCCCACCCATGGA
	SEQ ID No. 754:	5'- CCAGCTATCGATCATCGCCT
	SEQ ID No. 755:	5'- ACCAGCTATCGATCATGCC
5	SEQ ID No. 756:	5'- CAGCTATCGATCATGCCCTT
	SEQ ID No. 757:	5'- AGCTATCGATCATGCCTTG
	SEQ ID No. 758:	5'- GCTATCGATCATGCCCTGG
	SEQ ID No. 759:	5'- CTATCGATCATGCCCTTGGT
	SEQ ID No. 760:	5'- TTCGTGCGACTTGCATGTGT
10	SEQ ID No. 761:	5'- TCGATCATGCCCTTGGTAGG
	SEQ ID No. 762:	5'- ATCGATCATGCCCTTGGTAG
	SEQ ID No. 763:	5'- CACAGGCGACTTGCACCTT
	SEQ ID No. 764:	5'- CCACAGGCGACTTGCACCTT
	SEQ ID No. 765:	5'- TCCACAGGCGACTTGCACCT
15	SEQ ID No. 766:	5'- TCCTCCACAGGCGACTTGC
	SEQ ID No. 767:	5'- CCTCCACAGGCGACTTGC
	SEQ ID No. 768:	5'- CTCCACAGGCGACTTGC
	SEQ ID No. 769:	5'- ACAGGCGACTTGCACCTT
	SEQ ID No. 770:	5'- TCACCGGCTTAAGGTCAAAC
20	SEQ ID No. 771:	5'- CAACCCTCTCTCACACTCA
	SEQ ID No. 772:	5'- ACAACCCTCTCTCACACTCT
	SEQ ID No. 773:	5'- CCACAACCCTCTCTGACACT
	SEQ ID No. 774:	5'- AACCCCTCTCTCACACTCTAG
	SEQ ID No. 775:	5'- CACAACCCTCTCTCACACTC
25	SEQ ID No. 776:	5'- TCCACAACCCTCTCTCACAC
	SEQ ID No. 777:	5'- TTCCACAACCCTCTCTCAC
	SEQ ID No. 778:	5'- ACCCTCTCTCACACTCTAGT
	SEQ ID No. 779:	5'- GAGCCAGGTTGCCGCCTCG
	SEQ ID No. 780:	5'- AGGTCAAACCAACTCCCACATG
30	SEQ ID No. 781:	5'- ATGAGCCAGGTTGCCGCCTT

	SEQ ID No. 782:	5'- TGAGCCAGGTTGCCGCCTTC
	SEQ ID No. 783:	5'- AGGCTCCTCCACAGGCGACT
	SEQ ID No. 784:	5'- CAGGCTCCTCCACAGGCGAC
	SEQ ID No. 785:	5'- GCAGGCTCCTCCACAGGCGA
5	SEQ ID No. 786:	5'- TTCGCTCACCGGCTTAAGGT
	SEQ ID No. 787:	5'- GTTCGCTCACCGGCTTAAGG
	SEQ ID No. 788:	5'- GGTCGCTCACCGGCTTAAG
	SEQ ID No. 789:	5'- ATTCCACAACCCTCTCTCAC
	SEQ ID No. 790:	5'- TGACCCGACCGTGGTCGGCT
10	SEQ ID No. 791:	5'- CCCTCTCTCACACTCTAGTC
	SEQ ID No. 792:	5'- GAATTCCACAACCCTCTCTC
	SEQ ID No. 793:	5'- AGCCAGGTTGCCGCCTTCGC
	SEQ ID No. 794:	5'- GCCAGGTTGCCGCCTTCGCC
	SEQ ID No. 795:	5'- GGAATTCCACAACCCTCTCT
15	SEQ ID No. 796:	5'- GGAATTCCACAACCCTCTC
	SEQ ID No. 797:	5'- AACGCAGGCTCCTCCACAGG
	SEQ ID No. 798:	5'- CGGCTTAAGGTCAAACCAAC
	SEQ ID No. 799:	5'- CCGGCTTAAGGTCAAACCAA
	SEQ ID No. 800:	5'- CACCGGCTTAAGGTCAAACC
20	SEQ ID No. 801:	5'- ACCGGCTTAAGGTCAAACCA
	SEQ ID No. 802:	5'- ACCCAACATCCACACACAT
	SEQ ID No. 803:	5'- TCGCTGACCCGACCGTGGTC
	SEQ ID No. 804:	5'- CGCTGACCCGACCGTGGTCG
	SEQ ID No. 805:	5'- GACCCGACCGTGGTCGGCTG
25	SEQ ID No. 806:	5'- GCTGACCCGACCGTGGTCGG
	SEQ ID No. 807:	5'- CTGACCCGACCGTGGTCGGC
	SEQ ID No. 808:	5'- CAGGCGACTTGCGCCTTGA
	SEQ ID No. 809:	5'- TCATGCGGTATTAGCTCCAG
	SEQ ID No. 810:	5'- ACTAGCTAATCGAACGCAGG
30	SEQ ID No. 811:	5'- CATGCGGTATTAGCTCCAGT

	SEQ ID No. 812:	5'- CGCAGGCTCCTCCACAGGCG
	SEQ ID No. 813:	5'- ACGCAGGCTCCTCCACAGGC
	SEQ ID No. 814:	5'- CTCAGGTGTCATGCGGTATT
	SEQ ID No. 815:	5'- CGCCTTGACCCTCAGGTGT
5	SEQ ID No. 816:	5'- ACCCTCAGGTGTCATGCGGT
	SEQ ID No. 817:	5'- CCTCAGGTGTCATGCGGTAT
	SEQ ID No. 818:	5'- TTTGACCCTCAGGTGTCATG
	SEQ ID No. 819:	5'- GACCCTCAGGTGTCATGCGG
	SEQ ID No. 820:	5'- TGACCCTCAGGTGTCATGCG
10	SEQ ID No. 821:	5'- GCCTTGACCCTCAGGTGTC
	SEQ ID No. 822:	5'- TTGACCCTCAGGTGTCATGC
	SEQ ID No. 823:	5'- CCCTCAGGTGTCATGCGGTA
	SEQ ID No. 824:	5'- CCTTGACCCTCAGGTGTCA
	SEQ ID No. 825:	5'- CTTTGACCCTCAGGTGTCAT
15	SEQ ID No. 826:	5'- AGTTATCCCCCACCATGGA
	SEQ ID No. 827:	5'- CCAGCTATCGATCATCGCCT
	SEQ ID No. 828:	5'- ACCAGCTATCGATCATCGCC
	SEQ ID No. 829:	5'- CAGCTATCGATCATCGCCTT
	SEQ ID No. 830:	5'- AGCTATCGATCATCGCCTTG
20	SEQ ID No. 831:	5'- GCTATCGATCATCGCCTTGG
	SEQ ID No. 832:	5'- CTATCGATCATCGCCTTGGT
	SEQ ID No. 833:	5'- TTCGTGCGACTTGCATGTGT
	SEQ ID No. 834:	5'- TCGATCATCGCCTTGGTAGG
	SEQ ID No. 835:	5'- ATCGATCATCGCCTTGGTAG
25	SEQ ID No. 836:	5'- CACAGGCGACTTGCAGCCTT
	SEQ ID No. 837:	5'- CCACAGGCGACTTGCAGCCTT
	SEQ ID No. 838:	5'- TCCACAGGCGACTTGCAGCCT
	SEQ ID No. 839:	5'- TCCTCCACAGGCGACTTGCAG
	SEQ ID No. 840:	5'- CCTCCACAGGCGACTTGCAGC
30	SEQ ID No. 841:	5'- CTCCACAGGCGACTTGCAGCCTT

	SEQ ID No. 842:	5' - ACAGGCGACTTGCACCTTG
	SEQ ID No. 843:	5' - AGCCCCGGTTTCCCGGCGTT
	SEQ ID No. 844:	5' - CGCCTTCCTTTCCCTCCA
	SEQ ID No. 845:	5' - GCCCCGGTTTCCCGGCGTTA
5	SEQ ID No. 846:	5' - GCCGCCTTCCTTTCCCTC
	SEQ ID No. 847:	5' - TAGCCCCGGTTTCCCGGCGT
	SEQ ID No. 848:	5' - CCGGGTACCGTCAAGGCGCC
	SEQ ID No. 849:	5' - AAGCCGCCTTCCTTTCC
	SEQ ID No. 850:	5' - CCCGGTTTCCCGGCGTTAT
10	SEQ ID No. 851:	5' - CCGGC GTTATCCCAGTCTTA
	SEQ ID No. 852:	5' - AGCCGCCTTCCTTTCCCT
	SEQ ID No. 853:	5' - CCGCCTTCCTTTCCCTCC
	SEQ ID No. 854:	5' - TTAGCCCCGGTTTCCCGGCG
	SEQ ID No. 855:	5' - CCCGGCGTTATCCCAGTCTT
15	SEQ ID No. 856:	5' - GCCGGGTACCGTCAAGGCGC
	SEQ ID No. 857:	5' - GGCCGGGTACCGTCAAGGCG
	SEQ ID No. 858:	5' - TCCC GGCGTTATCCCAGTCT
	SEQ ID No. 859:	5' - TGGCCGGGTACCGTCAAGGC
	SEQ ID No. 860:	5' - GAAGCCGCCTTCCTTTTC
20	SEQ ID No. 861:	5' - CCCGGTTTCCCGGCGTTATC
	SEQ ID No. 862:	5' - CGGC GTTATCCCAGTCTTAC
	SEQ ID No. 863:	5' - GGCGTTATCCCAGTCTTACA
	SEQ ID No. 864:	5' - GCGTTATCCCAGTCTTACAG
	SEQ ID No. 865:	5' - CGGGTACCGTCAAGGCGCCG
25	SEQ ID No. 866:	5' - ATTAGCCCCGGTTTCCCGGC
	SEQ ID No. 867:	5' - AAGGGGAAGGCCCTGTCTCC
	SEQ ID No. 868:	5' - GGCCCTGTCTCCAGGGAGGT
	SEQ ID No. 869:	5' - AGGCCCTGTCTCCAGGGAGG
	SEQ ID No. 870:	5' - AAGGCCCTGTCTCCAGGGAGG
30	SEQ ID No. 871:	5' - GCCCTGTCTCCAGGGAGGT

	SEQ ID No. 872:	5'- CGTTATCCCAGTCTTACAGG
	SEQ ID No. 873:	5'- GGGTACCGTCAAGGCGCCGC
	SEQ ID No. 874:	5'- CGGCAACAGAGTTTACGAC
	SEQ ID No. 875:	5'- GGGGAAGGCCCTGTCTCCAG
5	SEQ ID No. 876:	5'- AGGGAAAGGCCCTGTCTCCA
	SEQ ID No. 877:	5'- GCAGCCGAAGCCGCCTTCC
	SEQ ID No. 878:	5'- TTCTTCCCCGGCAACAGAGT
	SEQ ID No. 879:	5'- CGGCACTTGTTCTTCCCCGG
	SEQ ID No. 880:	5'- GTTCTTCCCCGGCAACAGAG
10	SEQ ID No. 881:	5'- GGCACTTGTTCTTCCCCGGC
	SEQ ID No. 882:	5'- GCACTTGTTCTTCCCCGGCA
	SEQ ID No. 883:	5'- CACTTGTTCCTTCCCCGGCAA
	SEQ ID No. 884:	5'- TCTTCCCCGGCAACAGAGTT
	SEQ ID No. 885:	5'- TTGTTCTTCCCCGGCAACAG
15	SEQ ID No. 886:	5'- ACTTGTTCCTTCCCCGGCAAC
	SEQ ID No. 887:	5'- TGTTCTTCCCCGGCAACAGA
	SEQ ID No. 888:	5'- CTTGTTCTTCCCCGGCAACA
	SEQ ID No. 889:	5'- ACGGCACTTGTTCTTCCCCG
	SEQ ID No. 890:	5'- GTCCGCCGCTAACCTTTAA
20	SEQ ID No. 891:	5'- CTGGCCGGGTACCGTCAAGG
	SEQ ID No. 892:	5'- TCTGGCCGGGTACCGTCAAGG
	SEQ ID No. 893:	5'- TTCTGGCCGGGTACCGTCAA
	SEQ ID No. 894:	5'- CAATGCTGGCAACTAAGGTC
	SEQ ID No. 895:	5'- CGTCCGCCGCTAACCTTTA
25	SEQ ID No. 896:	5'- CGAAGCCGCCTTCCTTTT
	SEQ ID No. 897:	5'- CCGAAGCCGCCTTCCTTT
	SEQ ID No. 898:	5'- GCCGAAGCCGCCTTCCTTT
	SEQ ID No. 899:	5'- AGCCGAAGCCGCCTTCCTT
	SEQ ID No. 900:	5'- ACCGTCAAGGCCGCCCTG
30	SEQ ID No. 901:	5'- CCGTGGCTTCTGGCCGGT

	SEQ ID No. 902:	5'- GCTTTCTGGCCGGGTACCGT
	SEQ ID No. 903:	5'- GCCGTGGCTTCCTGGCCGGG
	SEQ ID No. 904:	5'- GGCTTCTGGCCGGGTACCG
	SEQ ID No. 905:	5'- CTTCTGGCCGGGTACCGTC
5	SEQ ID No. 906:	5'- TGGCTTCTGGCCGGGTACC
	SEQ ID No. 907:	5'- GTGGCTTCTGGCCGGGTAC
	SEQ ID No. 908:	5'- CGTGGCTTCTGGCCGGGTA
	SEQ ID No. 909:	5'- TTTCTGGCCGGGTACCGTCA
	SEQ ID No. 910:	5'- GGGAAAGGCCCTGTCTCCAGG
10	SEQ ID No. 911:	5'- CGAAGGGGAAGGCCCTGTCT
	SEQ ID No. 912:	5'- CCGAAGGGGAAGGCCCTGTC
	SEQ ID No. 913:	5'- GAAGGGGAAGGCCCTGTCTC
	SEQ ID No. 914:	5'- GGCGCCGCCCTGTTCGAACG
	SEQ ID No. 915:	5'- AGGCGCCGCCCTGTTCGAAC
15	SEQ ID No. 916:	5'- AAGGCGCCGCCCTGTTCGAA
	SEQ ID No. 917:	5'- CCCGGCAACAGAGTTTACG
	SEQ ID No. 918:	5'- CCCC GGCAACAGAGTTTAC
	SEQ ID No. 919:	5'- CCATCTGTAAGTGGCAGCCG
	SEQ ID No. 920:	5'- TCTGTAAGTGGCAGCCGAAG
20	SEQ ID No. 921:	5'- CTGTAAGTGGCAGCCGAAGC
	SEQ ID No. 922:	5'- CCCATCTGTAAGTGGCAGCC
	SEQ ID No. 923:	5'- TGTAAGTGGCAGCCGAAGCC
	SEQ ID No. 924:	5'- CATCTGTAAGTGGCAGCCGA
	SEQ ID No. 925:	5'- ATCTGTAAGTGGCAGCCGAA
25	SEQ ID No. 926:	5'- CAGCCGAAGCCGCCTTCCT
	SEQ ID No. 927:	5'- GGCAACAGAGTTTACGACC
	SEQ ID No. 928:	5'- CCGGCAACAGAGTTTACGA
	SEQ ID No. 929:	5'- TTCCCCGGCAACAGAGTTT
	SEQ ID No. 930:	5'- CTTCCCCGGCAACAGAGTTT
30	SEQ ID No. 931:	5'- TCCCCGGCAACAGAGTTTA

	SEQ ID No. 932:	5'- CCGTCCGCCGCTAACCTTT
	SEQ ID No. 933:	5'- CTTCCTCCGACTTACGCCGG
	SEQ ID No. 934:	5'- CCTCCGACTTACGCCGGCAG
	SEQ ID No. 935:	5'- TTCCTCCGACTTACGCCGGC
5	SEQ ID No. 936:	5'- TCCTCCGACTTACGCCGGCA
	SEQ ID No. 937:	5'- TCCGACTTACGCCGGCAGTC
	SEQ ID No. 938:	5'- CCGACTTACGCCGGCAGTCA
	SEQ ID No. 939:	5'- GCCTTCCTCCGACTTACGCC
	SEQ ID No. 940:	5'- CCTTCCTCCGACTTACGCCG
10	SEQ ID No. 941:	5'- GCTCTCCCCGAGCAACAGAG
	SEQ ID No. 942:	5'- CTCTCCCCGAGCAACAGAGC
	SEQ ID No. 943:	5'- CGCTCTCCCCGAGCAACAGA
	SEQ ID No. 944:	5'- CTCCGACTTACGCCGGCAGT
	SEQ ID No. 945:	5'- TCTCCCCGAGCAACAGAGCT
15	SEQ ID No. 946:	5'- CGACTTACGCCGGCAGTCAC
	SEQ ID No. 947:	5'- TCGGCACTGGGGTGTGTCCC
	SEQ ID No. 948:	5'- GGCACTGGGGTGTGTCCCC
	SEQ ID No. 949:	5'- CTGGGGTGTGTCCCCCAAC
	SEQ ID No. 950:	5'- CACTGGGGTGTGTCCCCCA
20	SEQ ID No. 951:	5'- ACTGGGGTGTGTCCCCCAA
	SEQ ID No. 952:	5'- GCACTGGGGTGTGTCCCCC
	SEQ ID No. 953:	5'- TGGGGTGTGTCCCCCAACA
	SEQ ID No. 954:	5'- CACTCCAGACTTGCTCGACC
	SEQ ID No. 955:	5'- TCACTCCAGACTTGCTCGAC
25	SEQ ID No. 956:	5'- CGGCACTGGGGTGTGTCCCC
	SEQ ID No. 957:	5'- CGCCTTCCTCCGACTTACGC
	SEQ ID No. 958:	5'- CTCCCCGAGCAACAGAGCTT
	SEQ ID No. 959:	5'- ACTCCAGACTTGCTCGACCG
	SEQ ID No. 960:	5'- CCCATGCCGCTCTCCCCGAG
30	SEQ ID No. 961:	5'- CCATGCCGCTCTCCCCGAGC

	SEQ ID No. 962:	5'- CCCCATGCCGCTCTCCCCGA
	SEQ ID No. 963:	5'- TCACTCGGTACCGTCTCGCA
	SEQ ID No. 964:	5'- CATGCCGCTCTCCCCGAGCA
	SEQ ID No. 965:	5'- ATGCCGCTCTCCCCGAGCAA
5	SEQ ID No. 966:	5'- TTCGGCACTGGGGTGTGTCC
	SEQ ID No. 967:	5'- TGCCGCTCTCCCCGAGCAAC
	SEQ ID No. 968:	5'- TTCACTCCAGACTTGCTCGA
	SEQ ID No. 969:	5'- CCCGCAAGAACAGATGCCTCCT
	SEQ ID No. 970:	5'- AGAACAGATGCCTCCTCGCGGG
10	SEQ ID No. 971:	5'- AAGAACAGATGCCTCCTCGCGG
	SEQ ID No. 972:	5'- CGCAAGAACAGATGCCTCCTCG
	SEQ ID No. 973:	5'- AAGATGCCTCCTCGCGGGCG
	SEQ ID No. 974:	5'- CCGCAAGAACAGATGCCTCCTC
	SEQ ID No. 975:	5'- GAAGATGCCTCCTCGCGGGC..
15	SEQ ID No. 976:	5'- CCCCCGCAAGAACAGATGCCTCC
	SEQ ID No. 977:	5'- CAAGAACAGATGCCTCCTCGCG
	SEQ ID No. 978:	5'- TCCTTCGGCACTGGGGTGTG
	SEQ ID No. 979:	5'- CCGCTCTCCCCGAGAACAG
	SEQ ID No. 980:	5'- TGCCTCCTCGCGGGCGTATC
20	SEQ ID No. 981:	5'- GACTTACGCCGGCAGTCACC
	SEQ ID No. 982:	5'- GGCTCCTCTCTCAGCGGGCCC
	SEQ ID No. 983:	5'- CCTTCGGCACTGGGGTGTGT
	SEQ ID No. 984:	5'- GGGGTGTGTCCCCCAACAC
	SEQ ID No. 985:	5'- GCCGCTCTCCCCGAGAACACA
25	SEQ ID No. 986:	5'- AGATGCCTCCTCGCGGGCGT
	SEQ ID No. 987:	5'- CACTCGGTACCGTCTCGCAT
	SEQ ID No. 988:	5'- CTCACTCGGTACCGTCTCGC
	SEQ ID No. 989:	5'- GCAAGAACAGATGCCTCCTCGC
	SEQ ID No. 990:	5'- CTCCAGACTTGCTCGACCGC
30	SEQ ID No. 991:	5'- TTACGCCGGCAGTCACCTGT

	SEQ ID No. 992:	5'- CTTCGGCACTGGGTGTGTC
	SEQ ID No. 993:	5'- CTCGCGGGCGTATCCGGCAT
	SEQ ID No. 994:	5'- GCCTCCTCGCGGGCGTATCC
	SEQ ID No. 995:	5'- ACTCGGTACCGTCTCGCATG
5	SEQ ID No. 996:	5'- GATGCCTCCTCGCGGGCGTA
	SEQ ID No. 997:	5'- GGGTGTGTCCCCCAACACC
	SEQ ID No. 998:	5'- ACTTACGCCGGCAGTCACCT
	SEQ ID No. 999:	5'- CTTACGCCGGCAGTCACCTG
	SEQ ID No. 1000:	5'- ATGCCTCCTCGCGGGCGTAT
10	SEQ ID No. 1001:	5'- GCGCCGCGGGCTCCTCTCTC
	SEQ ID No. 1002:	5'- GGTGTGTCCCCCAACACCT
	SEQ ID No. 1003:	5'- GTGTGTCCCCCAACACCTA
	SEQ ID No. 1004:	5'- CCTCGCGGGCGTATCCGGCA
	SEQ ID No. 1005:	5'- CCTCACTCGGTACCGTCTCG
15	SEQ ID No. 1006:	5'- TCCTCACTCGGTACCGTCTC
	SEQ ID No. 1007:	5'- TCGCGGGCGTATCCGGCATT
	SEQ ID No. 1008:	5'- TTTCACTCCAGACTTGCTCG
	SEQ ID No. 1009:	5'- TACGCCGGCAGTCACCTGTG
	SEQ ID No. 1010:	5'- TCCAGACTTGCTCGACCGCC
20	SEQ ID No. 1011:	5'- CTCGGTACCGTCTCGCATGG
	SEQ ID No. 1012:	5'- CGCGGGCGTATCCGGCATT
	SEQ ID No. 1013:	5'- GCGTATCCGGGATTAGGGCG
	SEQ ID No. 1014:	5'- GGGCTCCTCTCTCAGCGGCC
	SEQ ID No. 1015:	5'- TCCCCGAGCAACAGAGCTT
25	SEQ ID No. 1016:	5'- CCCCGAGCAACAGAGCTTA
	SEQ ID No. 1017:	5'- CCGAGCAACAGAGCTTACA
	SEQ ID No. 1018:	5'- CCATCCCATGGTTGAGCCAT
	SEQ ID No. 1019:	5'- GTGTCCCCCAACACCTAGC
	SEQ ID No. 1020:	5'- GCGGGCGTATCCGGCATTAG
30	SEQ ID No. 1021:	5'- CGAGCGGCTTTGGGTTTC

	SEQ ID No. 1022:	5'- CTTCACTCCAGACTTGCTC
	SEQ ID No. 1023:	5'- TTCCTTCGGCACTGGGGTGT
	SEQ ID No. 1024:	5'- CCGCCTTCCTCCGACTTACG
	SEQ ID No. 1025:	5'- CCCGCCTTCCTCCGACTTAC
5	SEQ ID No. 1026:	5'- CCTCCTCGCGGGCGTATCCG
	SEQ ID No. 1027:	5'- TCCTCGCGGGCGTATCCGGC
	SEQ ID No. 1028:	5'- CATTAGCGCCC GTTCCGG
	SEQ ID No. 1029:	5'- GCATTAGCGCCC GTTCCGG
	SEQ ID No. 1030:	5'- GGCATTAGCGCCC GTTCCG
10	SEQ ID No. 1031:	5'- GTCTCGCATGGGGCTTCCA
	SEQ ID No. 1032:	5'- GCCATGGACTTCACTCCAG
	SEQ ID No. 1033:	5'- CATGGACTTCACTCCAGAC
	SEQ ID No. 1037:	5'- ACCGTCTCAC AAGGAGCTT
	SEQ ID No. 1038:	5'- TACCGTCTCAC AAGGAGCTT
15	SEQ ID No. 1039:	5'- GTACCGTCTCAC AAGGAGCT
	SEQ ID No. 1040:	5'- GCCTACCCGTGTATTATCCG
	SEQ ID No. 1041:	5'- CCGTCTCAC AAGGAGCTTC
	SEQ ID No. 1042:	5'- CTACCCGTGTATTATCCGGC
	SEQ ID No. 1043:	5'- GGTACCGTCTCAC AAGGAGC
20	SEQ ID No. 1044:	5'- CGTCTCAC AAGGAGCTTCC
	SEQ ID No. 1045:	5'- TCTCAC AAGGAGCTTCCAC
	SEQ ID No. 1046:	5'- TACCCGTGTATTATCCGGCA
	SEQ ID No. 1047:	5'- GTCTCAC AAGGAGCTTCCA
	SEQ ID No. 1048:	5'- ACCCGTGTATTATCCGGCAT
25	SEQ ID No. 1049:	5'- CTCGGTACCGTCTCAC AAGG
	SEQ ID No. 1050:	5'- CGGTACCGTCTCAC AAGGAG
	SEQ ID No. 1051:	5'- ACTCGGTACCGTCTCAC AAG
	SEQ ID No. 1052:	5'- CGGCTGGCTCCATAACGGTT
	SEQ ID No. 1053:	5'- ACAAGTAGATGCCTACCCGT
30	SEQ ID No. 1054:	5'- TGGCTCCATAACGGTTACCT

	SEQ ID No. 1055:	5'- CAAGTAGATGCCTACCCGTG
	SEQ ID No. 1056:	5'- CACAAGTAGATGCCTACCCG
	SEQ ID No. 1057:	5'- GGCTCCATAACGGTTACCTC
	SEQ ID No. 1058:	5'- ACACAAGTAGATGCCTACCC
5	SEQ ID No. 1059:	5'- CTGGCTCCATAACGGTTACC
	SEQ ID No. 1060:	5'- GCTGGCTCCATAACGGTTAC
	SEQ ID No. 1061:	5'- GGCTGGCTCCATAACGGTTA
	SEQ ID No. 1062:	5'- GCTCCATAACGGTTACCTCA
	SEQ ID No. 1063:	5'- AAGTAGATGCCTACCCGTGT
10	SEQ ID No. 1064:	5'- CTCCATAACGGTTACCTCAC
	SEQ ID No. 1065:	5'- TGCCTACCCGTGTATTATCC
	SEQ ID No. 1066:	5'- TCGGTACCGTCTCACAAAGGA
	SEQ ID No. 1067:	5'- CTCACAAAGGAGCTTCCACT
	SEQ ID No. 1068:	5'- GTAGATGCCTACCCGTGTAT
15	SEQ ID No. 1069:	5'- CCTACCCGTGTATTATCCGG
	SEQ ID No. 1070:	5'- CACTCGGTACCGTCTCACAA
	SEQ ID No. 1071:	5'- CTCAGCGATGCAGTTGCATC
	SEQ ID No. 1072:	5'- AGTAGATGCCTACCCGTGTA
	SEQ ID No. 1073:	5'- GCGGCTGGCTCCATAACGGT
20	SEQ ID No. 1074:	5'- CCAAAGCAATCCAAGGTTG
	SEQ ID No. 1075:	5'- TCCATAACGGTTACCTCAC
	SEQ ID No. 1076:	5'- CCCGTGTATTATCCGGCATT
	SEQ ID No. 1077:	5'- TCTCAGCGATGCAGTTGCAT
	SEQ ID No. 1078:	5'- CCATAACGGTTACCTCACCG
25	SEQ ID No. 1079:	5'- TCAGCGATGCAGTTGCATCT
	SEQ ID No. 1080:	5'- GGCGGCTGGCTCCATAACGG
	SEQ ID No. 1081:	5'- AAGCAATCCAAGGTTGAGC
	SEQ ID No. 1082:	5'- TCACTCGGTACCGTCTCAC
	SEQ ID No. 1083:	5'- CCGAGTGTATTCCAGTCTG
30	SEQ ID No. 1084:	5'- CACAAGGAGCTTCCACTCT

	SEQ ID No. 1085:	5'- ACAAGGAGCTTCCACTCTC
	SEQ ID No. 1086:	5'- TCACAAGGAGCTTCCACTC
	SEQ ID No. 1087:	5'- CAGCGATGCAGTTGCATCTT
	SEQ ID No. 1088:	5'- CAAGGAGCTTCCACTCTCC
5	SEQ ID No. 1089:	5'- CCAGTCTGAAAGGCAGATTG
	SEQ ID No. 1090:	5'- CAGTCTGAAAGGCAGATTGC
	SEQ ID No. 1091:	5'- CGGCGGCTGGCTCCATAACG
	SEQ ID No. 1092:	5'- CCTCTCTCAGCGATGCAGTT
	SEQ ID No. 1093:	5'- CTCTCTCAGCGATGCAGTTG
10	SEQ ID No. 1094:	5'- TCTCTCAGCGATGCAGTTGC
	SEQ ID No. 1095:	5'- CTCTCAGCGATGCAGTTGCA
	SEQ ID No. 1096:	5'- CAATCCCAAGGTTGAGCCTT
	SEQ ID No. 1097:	5'- AATCCCAAGGTTGAGCCTTG
	SEQ ID No. 1098:	5'- AGCAATCCCAAGGTTGAGCC
15	SEQ ID No. 1099:	5'- CTCACTCGGTACCGTCTCAC
	SEQ ID No. 1100:	5'- GCAATCCCAAGGTTGAGCCT
	SEQ ID No. 1101:	5'- GCCTTGGACTTCACTTCAG
	SEQ ID No. 1102:	5'- CATAACGGTTACCTCACCGA
	SEQ ID No. 1103:	5'- CTCCTCTCTCAGCGATGCAG
20	SEQ ID No. 1104:	5'- TCGGCGGCTGGCTCCATAAC
	SEQ ID No. 1105:	5'- AGTCTGAAAGGCAGATTGCC
	SEQ ID No. 1106:	5'- TCCTCTCTCAGCGATGCAGT
	SEQ ID No. 1107:	5'- CCCAAGGTTGAGCCTTGGAC
	SEQ ID No. 1108:	5'- ATAACGGTTACCTCACCGAC
25	SEQ ID No. 1109:	5'- TCCAAGGTTGAGCCTTGGGA
	SEQ ID No. 1110:	5'- ATTATCCGGCATTAGCACCC
	SEQ ID No. 1111:	5'- CTACGTGCTGGTAACACAGA
	SEQ ID No. 1112:	5'- GCCGCTAGCCCCGAAGGGCT
	SEQ ID No. 1113:	5'- CTAGCCCCGAAGGGCTCGCT
30	SEQ ID No. 1114:	5'- CGCTAGCCCCGAAGGGCTCG

	SEQ ID No. 1115:	5'- AGCCCCGAAGGGCTCGCTCG
	SEQ ID No. 1116:	5'- CCGCTAGCCCCGAAGGGCTC
	SEQ ID No. 1117:	5'- TAGCCCCGAAGGGCTCGCTC
	SEQ ID No. 1118:	5'- GCTAGCCCCGAAGGGCTCGC
5	SEQ ID No. 1119:	5'- GCCCCGAAGGGCTCGCTCGA
	SEQ ID No. 1120:	5'- ATCCCAAGGTTGAGCCTTGG
	SEQ ID No. 1121:	5'- GAGCCTTGGACTTCACCTTC
	SEQ ID No. 1122:	5'- CAAGGTTGAGCCTTGGACTT
	SEQ ID No. 1123:	5'- GAGCTTCCACTCTCCTTGT
10	SEQ ID No. 1124:	5'- CCAAGGTTGAGCCTTGGACT
	SEQ ID No. 1125:	5'- CGGGCTCCTCTCTCAGCGAT
	SEQ ID No. 1126:	5'- GGAGCTTCCACTCTCCTTG
	SEQ ID No. 1127:	5'- GGGCTCCTCTCTCAGCGATG
	SEQ ID No. 1128:	5'- TCTCCTTGTGCGCTCTCCCCG
15	SEQ ID No. 1129:	5'- TCCTTGTGCGCTCTCCCCGAG
	SEQ ID No. 1130:	5'- AGCTTCCACTCTCCTTGTGTC
	SEQ ID No. 1131:	5'- CCACTCTCCTGTGCGCTCTC
	SEQ ID No. 1132:	5'- GGCTCCTCTCTCAGCGATGC
	SEQ ID No. 1133:	5'- CCTTGTGCGCTCTCCCCGAGC
20	SEQ ID No. 1134:	5'- CACTCTCCTTGTGCGCTCTCC
	SEQ ID No. 1135:	5'- ACTCTCCTTGTGCGCTCTCCC
	SEQ ID No. 1136:	5'- CTCTCCTTGTGCGCTCTCCCC
	SEQ ID No. 1137:	5'- GCGGGCTCCTCTCAGCGA
	SEQ ID No. 1138:	5'- GGCTCCATCATGGTTACCTC
25	SEQ ID No. 1142:	5'- CTTCCTCCGGCTTGCGCCGG
	SEQ ID No. 1143:	5'- CGCTCTCCGA(G/T)TGACTGA
	SEQ ID No. 1144:	5'- CCTCGGGCTCCTCCATC(A/T)GC

2. The method according to claim 1, wherein drink-spoiling microorganisms belonging to the genus *Zygosacchaeromyces* are detected with oligonucleotide probe SEQ ID No. 1.

5       3. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces bailii* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 5 to SEQ ID No. 21.

10       4. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces fermentati* is detected with oligonucleotide probe SEQ ID No. 22.

15       5. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces microellipsoides* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 23 to SEQ ID No. 24.

20       6. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces mellis* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 25 to SEQ ID No. 75.

25       7. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces rouxii* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 76 to SEQ ID No. 126.

8. The method according to claim 1, wherein the drink-spoiling microorganisms *Zygosacchaeromyces mellis* and *Zygosacchaeromyces rouxii* are detected simultaneously with oligonucleotide probe SEQ ID No. 127.

5        9. The method according to claim 1, wherein the drink-spoiling microorganism *Zygosacchaeromyces bisporus* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 128 to SEQ ID No. 142.

10        10. The method according to claim 1, wherein the drink-spoiling microorganism *Hanseniaspora uvarum* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 143 to SEQ ID No. 144.

15        11. The method according to claim 1, wherein the drink-spoiling microorganism *Candida intermedia* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 145 to SEQ ID No. 146.

20        12. The method according to claim 1, wherein the drink-spoiling microorganism *Candida parapsilosis* is detected with oligonucleotide probe SEQ ID No. 148.

13. The method according to claim 1, wherein the drink-spoiling microorganism *Candida crusei* (*Issatchenka orientalis*) is detected with oligonucleotide probe SEQ ID No. 149.

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14. The method according to claim 1, wherein the drink-spoiling microorganisms *Brettanomyces (Dekkera) anomala* and *Dekkera bruxellensis* are detected simultaneously with oligonucleotide probe SEQ ID No. 150.

15. The method according to claim 1, wherein the drink-spoiling microorganism *Brettanomyces (Dekkera) bruxellensis* is detected with oligonucleotide probe SEQ ID No. 151.

5        16. The method according to claim 1, wherein the drink-spoiling microorganism *Brettanomyces (Dekkera) naardenensis* is detected with oligonucleotide probe SEQ ID No. 152.

10        17. The method according to claim 1, wherein the drink-spoiling microorganism *Pichia membranaefaciens* is detected with oligonucleotide probe SEQ ID No. 153.

15        18. The method according to claim 1, wherein the drink-spoiling microorganisms *Pichia minuta* and *Pichia anomala* are detected simultaneously with oligonucleotide probe SEQ ID No. 154.

19. The method according to claim 1, wherein the drink-spoiling microorganism *Saccharomyces exiguum* is detected with oligonucleotide probe SEQ ID No. 157.

20        20. The method according to claim 1, wherein the drink-spoiling microorganism *Saccharomyces ludwigii* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 158 to SEQ ID No. 159.

25        21. The method according to claim 1, wherein the drink-spoiling microorganism *Saccharomyces cerevisiae* is detected with oligonucleotide probe SEQ ID No. 160.

22. The method according to claim 1, wherein the drink-spoiling microorganism *Mucor racemosus* is detected with oligonucleotide probe SEQ ID No. 163.

5        23. The method according to claim 1, wherein the drink-spoiling microorganism *Byssochlamys nivea* is detected with oligonucleotide probe SEQ ID No. 164.

10       24. The method according to claim 1, wherein the drink-spoiling microorganism *Neosartorya fischeri* is detected with oligonucleotide probe SEQ ID No. 165.

15       25. The method according to claim 1, wherein the drink-spoiling microorganisms *Aspergillus fumigatus* and *A. fischeri* are detected simultaneously with oligonucleotide probe SEQ ID No. 166.

20       26. The method according to claim 1, wherein the drink-spoiling microorganism *Talaromyces flavus* is detected with oligonucleotide probe SEQ ID No. 167.

25       27. The method according to claim 1, wherein the drink-spoiling microorganisms *Talaromyces bacillisporus* and *T. flavus* are detected simultaneously with oligonucleotide probe SEQ ID No. 168.

25       28. The method according to claim 1, wherein the drink-spoiling microorganism *Lactobacillus collinoides* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 169 to SEQ ID No. 269.

29. The method according to claim 1, wherein drink-spoiling microorganisms of the genus *Leuconostoc* are detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 270 to SEQ ID No. 271.

5        30. The method according to claim 1, wherein the drink-spoiling microorganisms *Leuconostoc mesenteroides* and *L. pseudomesenteroides* are detected simultaneously with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 272 to SEQ ID No. 301.

10        31. The method according to claim 1, wherein the drink-spoiling microorganism *Leuconostoc pseudomesenteroides* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 302 to SEQ ID No. 341.

15        32. The method according to claim 1, wherein the drink-spoiling microorganism *Oenococcus oenis* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 342 to SEQ ID No. 444.

20        33. The method according to claim 1, wherein drink-spoiling microorganisms of the genus *Weissella* are detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 445 to SEQ ID No. 495.

25        34. The method according to claim 1, wherein drink-spoiling microorganisms of the genus *Lactococcus* are detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 496 to SEQ ID No. 546.

30        35. The method according to claim 1, wherein drink-spoiling microorganisms of the genera *Acetobacter* and *Gluconobacter* are detected simultaneously with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 547 to SEQ ID No. 608.

36. The method according to claim 1, wherein drink-spoiling microorganisms of the genera *Acetobacter*, *Gluconobacter* and *Gluconoacetobacter* are detected simultaneously with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 609 to SEQ ID No. 842.

37. The method according to claim 1, wherein the drink-spoiling microorganism *Bacillus coagulans* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 843 to SEQ ID No. 932.

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38. The method according to claim 1, wherein drink-spoiling microorganisms of the genus *Alicyclobacillus* are detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 933 to SEQ ID No. 1033.

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39. The method according to claim 1, wherein the drink-spoiling microorganism *Alicyclobacillus acidoterrestris* is detected with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 1037 to SEQ ID No. 1138.

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40. The method according to claim 1, wherein the drink-spoiling microorganisms *Alicyclobacillus cycloheptanicus* and *A. herbarius* are detected simultaneously with at least one oligonucleotide probe selected from the group consisting of SEQ ID No. 1142 to SEQ ID No. 1144.

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41. The method according to claim 2, characterised in that the at least one oligonucleotide probe is used in combination with one or more competitor probes.

42. The method according to claim 41, characterised in that the oligonucleotide probe SEQ ID No. 1 is used in combination with one or more

competitor probes selected from the group consisting of SEQ ID No. 2 to SEQ ID No. 4.

43. The method according to claim 11, characterised in that the at least one  
5 oligonucleotide probe is used in combination with one or more competitor probes.

44. The method according to claim 43, characterised in that the  
oligonucleotide probe SEQ ID No. 146 is used in combination with competitor probe  
SEQ ID No. 147.

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45. The method according to claim 18, characterised in that the at least one  
oligonucleotide probe is used in combination with one or more competitor probes.

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46. The method according to claim 45, characterised in that the  
oligonucleotide probe SEQ ID No. 154 is used in combination with one or more  
competitor probes selected from the group consisting of SEQ ID No. 155 to SEQ ID  
No. 156.

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47. The method according to claim 21, characterised in that the at least one  
oligonucleotide probe is used in combination with one or more competitor probes.

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48. The method according to claim 47, characterised in that the  
oligonucleotide probe SEQ ID No. 160 is used in combination with one or more  
competitor probes selected from the group consisting of SEQ ID No. 161 to SEQ ID  
No. 162.

49. The method according to claim 38, characterised in that the at least one  
oligonucleotide probe is used in combination with one or more competitor probes.

50. The method according to claim 49, characterised in that the oligonucleotide probe SEQ ID No. 933 is used in combination with one or more competitor probes selected from the group consisting of SEQ ID No. 1034 to SEQ ID No. 1036.

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51. The method according to claim 39, characterised in that the at least one oligonucleotide probe is used in combination with one or more competitor probes.

10 52. The method according to claim 51, characterised in that the oligonucleotide probe SEQ ID No. 1044 is used in combination with the competitor probe SEQ ID No. 1139.

15 53. The method according to claim 51, characterised in that the oligonucleotide probe SEQ ID No. 1057 is used in combination with one or more competitor probes selected from the group consisting of SEQ ID No. 1140 to SEQ ID No. 1141.

54. The method according to any of claims 1 to 53, characterized in by comprising the following steps:

- 20     a) cultivating the drink-spoiling microorganisms contained in the sample,  
         b) fixing the drink-spoiling microorganisms contained in the sample,  
         c) incubating the fixed microorganisms with at least one oligonucleotide probe  
            optionally in combination with a competitor probe,  
         d) removing non-hybridised oligonucleotide probes,  
25     e) detecting and visualizing and optionally quantifying the drink-spoiling  
          microorganisms with the hybridized oligonucleotide probes.

55. The method according to any of claims 1 to 54, characterized in that the sample is a sample from non-alcoholic beverages.

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56. A kit for performing a method according to any of claims 1 to 55, containing at least one oligonucleotide according to claim 1.